

(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property Organization
International Bureau



(43) International Publication Date
23 January 2003 (23.01.2003)

PCT

(10) International Publication Number
WO 03/006103 A2

(51) International Patent Classification⁷: **A61N**

(21) International Application Number: PCT/US02/22161

(22) International Filing Date: 12 July 2002 (12.07.2002)

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Data:
60/304,955 12 July 2001 (12.07.2001) US

(71) Applicant (for all designated States except US): **MERCK & CO., INC.** [US/US]; Patent department, P.O. Box 2000 - RY60-30, Rahway, NJ 07065-0907 (US).

(72) Inventors; and

(75) Inventors/Applicants (for US only): **AUGUSTINE, Paul, R.** [US/US]; 8 Stima Avenue, Carteret, NJ 07008 (US). **BENNETT, Paul, B.** [US/US]; 3679 Hancock Lane, Doylestown, PA 18901 (US). **BUGIANESI, Randal, M.** [US/US]; 475 Milcrip Road, Bridgewater, NJ 09907 (US). **GARYANTES, Tina, A.** [US/US]; 18 Roberts Road, Warren, NJ 07059 (US). **IMREDY, John, P.** [US/US]; 861 Yorktown Street, Lansdale, PA 19446 (US). **KATH, Gary,**

S. [US/US]; 2671 Sky Top Drive, Scotch Plains, NJ 07076 (US). **MCMANUS, Owen, B.** [US/US]; 34 Robin Drive, Skillman, NJ 08558 (US).

(74) Agent: **VAN DYKE, Timothy, H.**; Van Dyke & Associates, P.A., 1630 Hillcrest Street, Orlando, FL 32803 (US).

(81) Designated States (*national*): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZM, ZW.

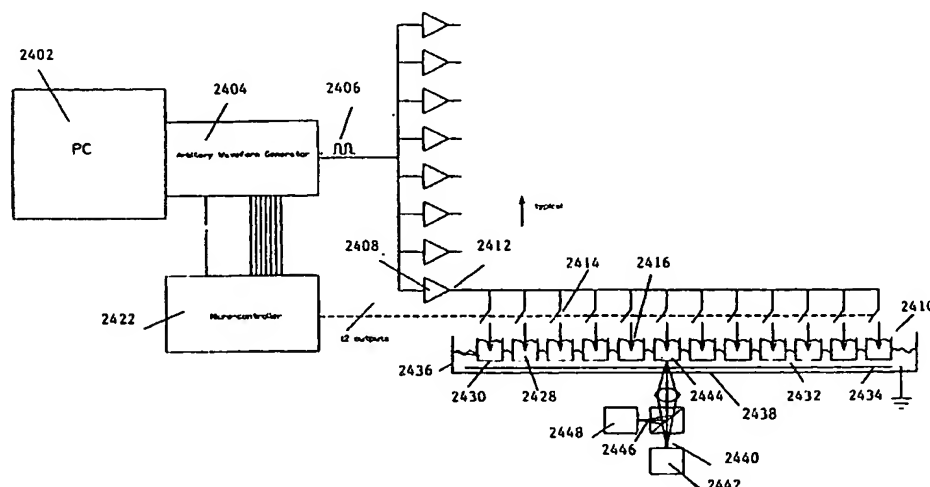
(84) Designated States (*regional*): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, SK, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG).

Published:

— without international search report and to be republished upon receipt of that report

[Continued on next page]

(54) Title: ELECTRICAL FIELD STIMULATION OF EUKARYOTIC CELLS



(57) Abstract: Methods of identifying activators and inhibitors of voltage-gated ion channels are provided in which the methods employ electrical field stimulation of the cells in order to manipulate the open/close state transition of the voltage-gated ion channels. This allows for more convenient, more precise experimental manipulation of these transitions, and, coupled with efficient methods of detecting the result of ion flux through the channels, provides methods that are especially suitable for high throughput screening.

WO 03/006103 A2

BEST AVAILABLE COPY



For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

TITLE OF THE INVENTION

ELECTRICAL FIELD STIMULATION OF EUKARYOTIC CELLS

CROSS-REFERENCE TO RELATED APPLICATIONS

5 The subject application is related to co-pending provisional application
no. 60/304,955, filed July 12, 2001, to which priority is claimed under 35 USC §
119(e).

STATEMENT REGARDING FEDERALLY-SPONSORED R&D

10 Not applicable.

REFERENCE TO MICROFICHE APPENDIX

Not applicable.

15 FIELD OF THE INVENTION

 The present invention is directed to methods and associated
apparatuses for stimulating eukaryotic cells by the application of electric fields. The
electric fields are produced by certain arrangements of electrodes that create an
electric potential difference in the environment of the cells, resulting in a change in
20 membrane potential of the cells. The change in membrane potential affects various
physiological processes within the cells, including the opening and closing of voltage-
gated ion channels. The ability to alter the open/close transitions of voltage-gated ion
channels by the application of electric fields as described herein provides for novel
methods of screening compounds for the ability to modulate the activity of voltage-
25 gated ion channels.

BACKGROUND OF THE INVENTION

 Certain molecular events in eukaryotic cells depend on the existence or
magnitude of an electric potential gradient across the plasma (*i.e.*, outer) membrane of
30 the cells. Among the more important of such events is the movement of ions across
the plasma membrane through voltage-gated ion channels. Voltage-gated ion
channels form transmembrane pores that open in response to changes in cell
membrane potential and allow ions to pass through the membrane. Voltage-gated ion
channels have many physiological roles. They have been shown to be involved in

maintaining cell membrane potentials and controlling the repolarization of action potentials in many types of cells (Bennett et al., 1993, Cardiovascular Drugs & Therapy 7:195-202; Johnson et al., 1999, J. Gen. Physiol. 113:565-580; Bennett & Shin, "Biophysics of voltage-gated sodium channels," in Cardiac Electrophysiology: From Cell to Bedside, 3rd edition, D. Zipes & J. Jalife, eds., 2000, W.B. Saunders Co., pp.67-86; Bennett & Johnson, "Molecular physiology of cardiac ion channels," Chapter 2 in Basic Cardiac Electrophysiology and Pharmacology, 1st edition, A. Zasa & M. Rosen, eds., 2000, Harwood Academic Press, pp. 29-57). Moreover, mutations in sodium, calcium, or potassium voltage-gated ion channel genes leading to defective channel proteins have been implicated in a variety of disorders including the congenital long QT syndromes, ataxia, migraine, muscle paralysis, deafness, seizures, and cardiac conduction diseases, to name a few (Bennett et al., 1995, Nature 376:683-685; Roden et al., 1995, J. Cardiovasc. Electrophysiol. 6:1023-1031; Kors et al., 1999, Curr. Opin. Neurol. 12:249-254; Lehmann et al., 1999, Physiol. Rev. 79:1317-1372; Holbauer & Heufelder, 1997, Eur. J. Endocrinol. 136:588-589; Naccarelli & Antzelevitch, 2000, Am. J. Med. 110:573-581).

Several types of voltage-gated ion channels exist. Voltage-gated potassium channels establish the resting membrane potential and modulate the frequency and duration of action potentials in neurons, muscle cells, and secretory cells. Following depolarization of the membrane potential, voltage-gated potassium channels open, allowing potassium efflux and thus membrane repolarization. This behavior has made voltage-gated potassium channels important targets for drug discovery in connection with a variety of diseases. Dysfunctional voltage-gated potassium channels have been implicated in a number of diseases and disorders. Wang et al., 1998, Science 282:1890-1893 have shown that the voltage-gated potassium channels KCNQ2 and KCNQ3 form a heteromeric potassium ion channel known as the "M-channel." Mutations in KCNQ2 and KCNQ3 in the M-channel are responsible for causing epilepsy (Biervert et al., 1998, Science 279:403-406; Singh et al., 1998, Nature Genet. 18:25-29; Schroeder et al., Nature 1998, 396:687-690).

Voltage-gated sodium channels are transmembrane proteins that are essential for the generation of action potentials in excitable cells (Catterall, 1993, Trends Neurosci. 16:500-506). In mammals, voltage-gated sodium channels consist of a macromolecular assembly of α and β subunits with the α subunit being the pore-forming component. α subunits are encoded by a large family of related genes, with

some α subunits being present in the central nervous system (Noda et al., 1986, Nature 322:826-828; Auld et al., 1988, Neuron 1:449-461; Kayano et al., 1988, FEBS Lett. 228:187-194) and others in muscle (Rogart et al., 1989, Proc. Natl. Acad. Sci. USA 86:8170-8174; Trimmer et al., 1989, Neuron 3:33-49).

5 Voltage-gated calcium channels are transmembrane proteins that in the open configuration allow the passive flux of Ca^{2+} ions across the plasma membrane, down the electrochemical gradient. They mediate various cell functions, including excitation-contraction coupling, signal transduction, and neurotransmitter release.

Current methods of drug discovery often involve assessing the
10 biological activity (*i.e.*, screening) of tens or hundreds of thousands of compounds in order to identify a small number of those compounds having a desired activity. In many high throughput screening programs, it is desirable to test as many as 50,000 to 100,000 compounds per day. Unfortunately, current methods of assaying the activity of voltage-gated ion channels are ill suited to the needs of a high throughput screening
15 program. Current methods often rely on electrophysiological techniques. Standard electrophysiological techniques involve "patching" or sealing against the cell membrane with a glass pipette followed by suction on the glass pipette, leading to rupture of the membrane patch (Hamill et al., 1981, Pflugers Arch. 391:85-100). This has limitations and disadvantages. Accessing the cell interior may alter the cell's
20 response properties. The high precision optical apparatuses necessary for micromanipulating the cells and the pipettes make simultaneous recording from more than a few cells at a time impossible. Given these difficulties, the throughput that can be achieved with electrophysiological techniques falls far short of that necessary for high throughput screening.

25 Various techniques have been developed as alternatives to standard methods of electrophysiology. For example, radioactive flux assays have been used in which cells are loaded with a radioactive tracer (*e.g.*, $^{86}\text{Rb}^+$, $^{22}\text{Na}^+$, $[^{14}\text{C}]$ -guanidinium) and the efflux of the dye is monitored. Cells loaded with the tracer are exposed to compounds and those compounds that either enhance or diminish the
30 efflux of the tracer are identified as possible activators or inhibitors of ion channels in the cells' membranes.

Assays that measure the change in a cell's membrane potential due to the change in activity of an ion channel have been developed. Such assays often employ voltage sensitive dyes that redistribute between the extracellular environment

and the cell's interior based upon a change in membrane potential and that have a different fluorescence spectrum depending on whether they are inside or outside the cell. A related assay method uses a pair of fluorescent dyes capable of fluorescence resonance energy transfer to sense changes in membrane potential. For a description of this technique, see González & Tsien, 1997, *Chemistry & Biology* 4:269-277. See also González & Tsien, 1995, *Biophys. J.* 69:1272-1280 and U.S. Patent No. 5,661,035. Other methods employ ion selective indicators such as calcium dependent fluorescent dyes to monitor changes in Ca^{2+} influx during opening and closing of calcium channels.

Ideally, methods of screening against voltage-gated ion channels require that the transmembrane potential of the cells being assayed be controlled and/or that the ion channels studied be cycled between open and closed states. This has been done in various ways. In standard electrophysiological techniques, the experimental set-up allows for direct manipulation of membrane potential by the voltage clamp method (Hodgkin & Huxley, 1952, *J. Physiol. (Lond.)* 153:449-544), *e.g.*, changing the applied voltage or injecting various ions into the cell. In other methods, changing the extracellular K^{+} concentration from a low value (*e.g.*, 5 mM) to a higher value (*e.g.*, 70-80 mM) results in a change in the electrochemical potential for K^{+} due to the change in the relative proportion of intracellular and extracellular potassium. This results in a change in the transmembrane electrical potential towards a more depolarized state. This depolarization can activate many voltage-gated ion channels, *e.g.*, voltage-gated calcium, sodium, or potassium channels. Alternatively, Na^{+} channels can be induced into an open conformation by the use of toxins such as veratridine or scorpion venom (Strichartz et al., 1987, *Ann. Rev. Neurosci.* 10:237-267; Narahashi & Harman, 1992, *Meth. Enzymol.* 207:620-643). While sometimes effective, such experimental manipulations may alter the channel pharmacology, can be awkward to perform, and can lead to artifactual disturbances in the system being studied.

Electrical field stimulation of cells has been performed on a single cell by sealing a glass microelectrode to the cell membrane. Rupture of the sealed patch of cell membrane resulted in an electrical connection between the interior fluid in the glass microelectrode and the fluid within the cell that was used to stimulate the cell via an electronic pulse generator. The electrophysiological response of the cell was measured via a sensitive electronic amplifier. The disadvantage of this technique is

that only one cell at a time was tested and it is a tedious and time consuming operation to seal the microelectrode to an individual cell.

HEK293 cells have been grown on a silicon chip made up of an array of field-effect transistors. Some of the cells were positioned over the gate region of the transistors, thus having portions of their plasma membranes overlying the source and the drain. When a patch pipette in such cells manipulated the intracellular voltage, Maxi-K potassium channels in the cells' plasma membranes were opened. This led to current flow in the region between the cells' membrane and the transistor. This current flow modulated the source-drain current, which could be detected by an appropriate device. The chip plus cells was said to have potential as a sensor and as a prototype for neuroprosthetic devices. See Straub et al., 2001, Nature Biotechnol. 19:121-124; Neher, 2001, Nature Biotechnol. 19:114.

SUMMARY OF THE INVENTION

The present invention is directed to methods of identifying activators and inhibitors of voltage-gated ion channels in which the methods employ electrical field stimulation of the cells via extracellular electrodes in order to manipulate the open/close state transitions of the voltage-gated ion channels. This allows for more convenient, more precise manipulation of these transitions, and, coupled with efficient methods of detecting ion flux or membrane potential, results in methods that are especially suitable for high throughput screening in order to identify substances that are activators or inhibitors of voltage-gated ion channels.

The present invention also provides apparatuses for use in the above-described methods. In particular, modifications of standard multiwell tissue culture plates are provided where the modified multiwell tissue culture plates have electrodes that can alter the transmembrane electric potential of cells in the wells of the plates, thus altering the ratio of open/close states of voltage-gated ion channels in the cells.

BRIEF DESCRIPTION OF THE DRAWINGS

Figure 1A shows a top view of one embodiment of the present invention. This embodiment comprises a glass slide 1 in which or upon which are a gold positive electrode 2 and a gold negative electrode 3 spaced such that a gap 4 of about 25 μm to 100 μm exists between the electrodes. The electrodes together with spacers 5 (here shown as plastic strips) arranged generally at right angles to the

electrodes define a series of wells 6 about 100 μm deep into which cells can be placed and/or grown. Figure 1B shows a cross-sectional side view of the embodiment of Figure 1A. In this embodiment, the identities of the positive and negative electrodes can be interchanged, if desired. The electrodes need not be made from gold; other
5 conductive materials may be used. Also, the spacers need not be plastic; other non-conductive materials may be used.

Figure 2A shows a top view of an embodiment of the present invention in which a typical 96 well plate contains electrodes within each well. Figure 2B shows a cross-sectional side view of one of the wells in Figure 2A. The well has a
10 first electrode 1 (here shown as a positive electrode) on the side 2 of the well, a second electrode 3 (here shown as a negative electrode) on the bottom 4 of the well, a strip of an optional insulating material 5 on the bottom of the well, and a cell 6 at the bottom of the well. A single cell is shown merely for convenience of illustration; in most cases a plurality of cells would be in the bottom of the well. The sides 2 of the
15 well are made of a non-conducting material such as plastic and the bottom of the well is made from a conducting material such as indium tin oxide (ITO). The well is shown with a fluid level 7 sufficient to completely cover the cell 6 and the second electrode 3 at the bottom 4 of the well and to reach the first electrode 1 on the side 2 of the well. The well is not drawn to scale with respect to Figure 2A. Figure 2C
20 shows an alternative arrangement of electrodes in a well. In this embodiment, both the positive electrode 1 and the negative electrode 2 are in the bottom 3 of the well. In this embodiment, the sides 4 and bottom 3 of the well are made of non-conducting material such as plastic. The fluid level 5 is such as to cover the cells 6 as well as the positive 1 and negative 2 electrodes.

25 Figure 3 shows a single well 1 from an embodiment of the invention where first 2 and second 3 electrodes are interdigitating and have been chemically etched on a layer of conductive material on the surface of a glass substrate 4. The well is generally circular with a 3 mm diameter. The electrodes are 10 μm wide and have a spacing of 160 μm . Either the first 2 or the second 3 electrodes may function
30 as the positive electrode. The width of the electrodes and the spacing between the electrodes can be varied. The width is preferably between 1 and 10 μm ; the spacing between the electrodes is preferably 5 μm to 160 μm . In particularly preferred embodiments, the spacing between the electrodes is at least as great as a typical diameter of a eukaryotic cell (*i.e.*, about 40 μm to 50 μm).

Figure 4A and 4B illustrates an embodiment in which wells are formed by attaching a well frame onto the substrate. Figure 4A shows an exploded view of the embodiment containing a well frame 1 the openings 2 of which form the wells on the substrate 3 where the well frame 1 is attached to the substrate 3 (*e.g.*, by gluing it in place), a contact guide plate 5 with a spring loaded contact 6, and a printed circuit board (PCB) 7. The substrate holder 4 is used to hold the assembled device in position on a measuring instrument such as a microscope or fluorescent plate reader (not shown). The PCB 7 contains connections through which the electrodes (not shown) can be linked to a pulse generator (not shown). Figure 4B shows an assembled view.

Figure 5 shows an arrangement of interdigitating electrodes formed upon a substrate that contains virtual wells. Virtual wells are described further herein.

Figure 6 shows a single well from an embodiment of the invention where two substantially parallel plates 1 have their opposing surfaces coated with conductive layers 2 between which is sandwiched a droplet of fluid containing the cells to be tested 3. One conductive layer is a positive electrode (here the upper conductive layer 4) while the other conductive layer is a negative electrode (here the lower conductive layer 5). Of course, the identity of the electrodes could be reversed, with the upper conductive layer being the negative electrode and the lower conductive layer being the positive electrode). In particular versions of this embodiment, the plates are glass and the conductive layer is indium tin oxide (ITO). The conductive layer preferably has a thickness of about 200 Å to 2,000 Å, or 500 Å to 1,500 Å, or 800 Å to 1,200 Å.

Figure 7 shows a single well 3 from an embodiment of the invention where one of the electrodes is a thin coating of conductive material 2 on the surface of a flat substrate 1 and forms the bottom 10 of the well. The other electrode 7 enters the well 3 from above and makes contact with the fluid 5 within the well 3. Electrode 7 is shown in cut-away view. Electrode 7 contains a central conductive material portion 8 that is surrounded by an insulator 6. For the sake of simplicity, a single cell 4 is shown in the well. Generally, at least 10^5 cells would be present in the well. The conductive layer preferably has a thickness of about 200 Å to 2,000 Å, or 500 Å to 1,500 Å, or 800 Å to 1,200 Å.

Figure 8 shows a single well 4 from an embodiment of the invention where the bottom of the well 4 is a filter membrane 12 upon which cells can be

grown. For simplicity, a single cell 8 is shown on the filter membrane 12. The well 4 is located in a trough 2 having a glass bottom 1 and filled with a first fluid 3. One electrode 7 enters the well 4 from above and makes contact with a second fluid 5 within the well 4. Electrode 7 contains a central conductive material portion that is surrounded by an insulator 6 and is connected to a pulse generator (not shown) by a first lead 9. A second electrode 11 is positioned within the first fluid 3 and is connected to the pulse generator by a second lead 10. The second electrode 11 is shown in cut-away view. The second electrode 11 actually forms a circle in the bottom of the well 4. Either the first electrode 7 is the positive electrode while the second electrode 11 is the negative electrode or the first electrode 7 is the negative electrode while the second electrode 11 is the positive electrode.

Figure 9A shows a single well 2 from an embodiment of the invention where both the positive 5 and negative 8 electrodes enter the well 2 from above. The well 2 contains fluid 3 in which a single cell 9 is shown, although generally a plurality of cells will be present in the well 2. The positive electrode 5 is connected to a pulse generator (not shown) by a positive lead 6. The negative electrode 8 is connected to the pulse generator by a negative lead 7. Both electrodes are embedded in an insulator 4. The positive 5 and negative 8 electrodes traverse the interior of the insulator 4 such that the positive 5 and negative 8 electrodes are generally perpendicular to a glass plate 1 that forms the bottom of the well 2. However, when the positive 5 and negative 8 electrodes exit the bottom 10 of the insulator 4, the positive 5 and negative 8 electrodes are each bent into a 90° angle so that they lie on and parallel to the bottom 10 of the insulator 4. Figure 9B is a view looking up from the glass plate 1 that forms the bottom of the well 2 and shows the arrangement of the bent portion of the positive 5 and negative 8 electrodes lying on bottom of the insulator 4.

Figure 10A shows an embodiment of the invention where both the positive 5 and negative 8 electrodes enter the well 2 from above and the positive 5 and negative 8 electrodes are arranged in a manner similar to that of a co-axial cable. The positive electrode 5 is embedded in an insulator 4 with the negative electrode 8 coating the outside of the insulator 4. The positive electrode 5 is connected to a pulse generator (not shown) by a positive lead 6. The negative electrode 8 is connected to the pulse generator by a negative lead 7. The well 2 contains fluid 3 in which a single cell 9 is shown, although generally a plurality of cells will be present in the well 2. A

glass plate 1 forms the bottom of the well 2. Figure 10B shows a view looking up from below the positive 5 and negative 8 electrodes.

Figure 11 shows an embodiment of the invention similar to the embodiment shown in Figure 8 except that in Figure 11 the electrode 7 that enters the well from above is not surrounded by an insulator but instead is within a pipette tip 6 and makes contact with a first fluid 5 also within the pipette tip 6 that is co-extensive with the first fluid 5 in the well 4. This arrangement has the advantage of minimizing the formation of bubbles in the first fluid 5 in the area at the end of the electrode 7. The bottom of the well 4 is a filter membrane 12 upon which cells can be grown. For simplicity, a single cell 8 is shown on the filter membrane 12. The well 4 sits in a trough 2 having a glass bottom 1 and filled with a second fluid 3. Electrode 7 is connected to a pulse generator (not shown) by a first lead 9. A second electrode 11 is positioned within the second fluid 3 and is connected to the pulse generator by a second lead 10. The second electrode 11 is shown in cut-away view. The second electrode 11 actually forms a circle in the bottom of the well 4. Either electrode can be the positive or negative electrode.

Figure 12A-B shows an embodiment that is similar to the embodiment of Figure 7 in having one electrode enter from above while the other electrode forms the bottom of the wells. Figure 12A is a side cross-sectional view that shows a substrate that is a 96-well microtiter plate in which one electrode 1 is a layer of a conductive material such as ITO that forms the bottom of the wells 2. The other electrode 3 enters the wells from above and makes contact with the fluid in the wells (fluid not shown). The electrodes are connected to an electrical pulse generator 4 by leads 5. Either electrode may be the positive or negative electrode. An alternative embodiment, similar to that shown, is to replace the bottom of standard 96, 384, 1536, or 3456 well plates with a conductive material such as ITO, which forms one electrode. The second electrode is lowered into each well from above. Contact to the ITO electrode can be made via electrically conducting silver epoxide or by placing a 3 M KCl (or similar salt solution) in alternate wells as the contact to the ITO bottoms from a platinum wire. Figure 12B shows a top view of the substrate.

Figure 13A-B shows an embodiment comprising two multiwell substrates containing virtual wells. Figure 13A is a side cross-sectional view that shows the top substrate 1 approaching the bottom substrate 2. The top electrode 3 is made of a conducting material such as ITO and forms the bottom of the virtual wells 4

of the top substrate 1. Similarly, the bottom electrode 5 is made of a conducting material such as ITO and forms the bottom of the virtual wells 6 of the bottom substrate 2. A thin layer of TEFLON® or a similar hydrophobic material 11 covers the surfaces of the conducting material on the substrates. Circular areas of the surface of the substrate that lack TEFLON® are relatively hydrophilic and form the virtual wells. The TEFLON® layer is about 0.5 μm to 100 μm thick. The top 3 and bottom 5 electrodes are connected to an electrical pulse generator 6 by leads 7. The left most wells of the apparatus are shown containing fluid drops. The top drop 8 might contain a substance such as a drug or a compound to be tested while the bottom drop 9 might contain cells expressing a voltage-gated ion channel. Figure 13B shows the apparatus after the top 1 and bottom 2 substrates have moved close enough together so that the top 8 and bottom 9 drops have mixed. 10 is a spacer (not shown in Figure 13A) that helps to align the top 1 and bottom 2 substrates and keeps the substrates the proper distance apart for mixing of the drops.

Figure 14 illustrates the principles of electrical field stimulation of cells.

Figure 15 shows two wells from an embodiment where one electrode enters the wells from above 1 while the second electrode is formed from the transparent ITO-coated bottom 2 of the transparent substrate 3 that is in contact with a highly conductive metal grounding grid 4. The dashed lines with arrowheads illustrate how current flows from the electrodes that enter from above 1 through a buffered salt solution 5 and the cells 6 and through the ITO layer 2 and the metal grounding grid 4. Arrows 7 within the substrate 3 illustrate how light from a source used in the detection system (not shown) would pass in the upward direction through the transparent substrate 3 and the ITO layer 2 into the cells 6 and then be re-emitted by the cells 6 as fluorescence and pass downward to a detector (not shown). Optional adhesive seals 8 that can be used to attach the wells to the ITO-coated substrate 3 are shown. The thickness of the ITO layer is preferably about 200 Å to 2,000 Å, or 500 Å to 1,500 Å, or 800 Å to 1,200 Å.

Figure 16A shows two wells of a multiwell embodiment having a conductive layer 1 such as ITO that forms the bottom of the wells. The positive electrode 2 enters the left well 3 from above while the negative electrode 4 enters the right well 5 from above. The transparent layer of a conductive material 1 such as ITO coats a transparent substrate 7 such as glass. The dotted line with an arrowhead

shows the path of current flow. Of course, the identity of the positive and negative electrodes could be reversed. Cells 8 are shown in fluid 9 within the wells. Optional adhesive seals 10 that can be used to attach the wells to the ITO-coated substrate 7 are shown. Light path is indicated by arrows in the substrate. Figure 16B shows a side cut-away view of this embodiment that illustrates how the positive 2 and negative 4 electrodes might be connected to a pulse generator 11. Also shown is the transparent conductive layer 6 coating the transparent substrate 7. Figure 16C shows a top view of the embodiment that illustrates the alternating pattern of positive and negative electrodes. Figure 16D is a photograph of this embodiment that has been partially disassembled. The wells are formed by a well frame 12 that is attached to the glass substrate 13 that is has been coated with ITO. During normal operation, the substrate will cover all the wells. For the purpose of illustration, this view shows only part of the substrate.

Figure 17 shows a graphical representation of data obtained from an embodiment of the invention similar to that depicted in Figure 16. The data represent Ca^{2+} influx into HEK293 cells that have been transfected to express the human $\alpha 1\text{H}$ T-type voltage-gated calcium channel (GenBank accession no. AF073931). Ca^{2+} influx occurred when the T-type channels opened and was measured by detecting fluorescent emission at 520-560 nm of the calcium indicator dye Fluo4 that had been excited at 480 nm. At the time points indicated, a preselected voltage was applied through the electrodes. This resulted in the opening of a portion of the T-type channels, allowing Ca^{2+} influx. This caused a spike in the fluorescent emission at 520-560 nm by the calcium indicator dye Fluo4. The spike gradually decayed, as shown.

Figure 18A-B shows a nucleotide sequence encoding the human PN3 sodium channel (SEQ.ID.NO.:1). Figure 18C shows the corresponding amino acid sequence (SEQ.ID.NO.:2). From GenBank accession no. AF117907.

Figure 19A-B shows a nucleotide sequence encoding the $\alpha 1\text{H}$ subunit of the human T-type calcium channel (SEQ.ID.NO.:3). Figure 19C shows the corresponding amino acid sequence (SEQ.ID.NO.:4). From GenBank accession no. AF073931.

Figure 20A-B shows a nucleotide sequence encoding a splice variant of the $\alpha 1\text{B}$ subunit of the human N-type calcium channel (SEQ.ID.NO.:5). Figure

20C shows the corresponding amino acid sequence (SEQ.ID.NO.:6). From GenBank accession no. M94172.

Figure 21A-B shows a nucleotide sequence encoding a splice variant of the $\alpha 1B$ subunit of the human N-type calcium channel (SEQ.ID.NO.:7). Figure 21C shows the corresponding amino acid sequence (SEQ.ID.NO.:8). From GenBank accession no. M94173.

Figure 22A-B shows a nucleotide sequence encoding the human calcium channel $\alpha 1A$ isoform 1A-1 subunit (SEQ.ID.NO.:9). Figure 22C shows the corresponding amino acid sequence (SEQ.ID.NO.:10). From GenBank accession no. AF004884.

Figure 23A-B shows a nucleotide sequence encoding the human calcium channel $\alpha 1A$ isoform 1A-2 subunit (SEQ.ID.NO.:11). Figure 23C shows the corresponding amino acid sequence (SEQ.ID.NO.:12). From GenBank accession no. AF004883.

Figure 24 shows a schematic diagram of one embodiment of a EFS system utilizing a computer, voltage generator, amplifier, membrane bottom wells, common trough, and fluorescence detector, *inter alia*.

Figure 25 is a photograph showing an electrode head embodiment especially adapted for use with a 96 well tray.

Figure 26 is a photograph showing a trough embodiment for use in conjunction with the electrode head embodiment shown in Figure 25.

Figure 27 is a photograph showing the trough embodiment of Figure 26 with a multi-screen well tray positioned therein.

Figure 28 is a photograph showing the assembled electrode head, trough and multiscreen.

Figure 29 shows a graphical representation of data obtained from an embodiment of the invention similar to that depicted in Figure 28. The data represent a membrane potential change in HEK293 cells that have been transfected to express

human PN1 voltage-gated sodium channel. Each plot represents a row (12wells) A-H of a 96-well plate. Each column of the 96-well plate data was acquired for 15 seconds on a VIPR™. Stimulation pulse protocol was applied during the data acquisition as follows; 2s baseline was followed with a 2ms square pulse, Amplitude = 20mA,
5 Frequency = 10 Hz, Duration = 5s.

Figure 30 is a bar graph representation of the peak ration change of data depicted in Figure 29. 1 μ M TTX a specific and potent blocker of tetrodotoxin (TTX)-sensitive voltage-gated sodium channels is present in wells E1, F1, G1, H1,
10 A12, B12, C12 and D12. In addition well A11 contains an internal standard for blocking TTX-sensitive voltage-gated sodium channels. Z-score is a measure of the difference in the uninhibited and inhibited signal divided by the sum of the standard deviations.

15 Figure 31 shows the effects of increasing concentrations of TTX (upper panel) and of Compound A (lower panel) on the EFS-stimulated depolarization signal in HEK293/PN1 cells. The IC_{50} s obtained in these experiments are comparable to those obtained through other techniques. The high Hill coefficients, nH , result from the threshold nature of the stimulation protocol.

20 Figure 32 is a photograph showing an alternative embodiment. Figure 32 shows an electrode head similar to that shown in Figure 25, and a copper electrode plate. This embodiment is especially adapted for use with Caco-2 multiscreens (Millipore, Bedford, MA).

25 Figure 33 is a photograph similar to that shown in Figure 32 except that the copper electrode plate has been turned over to show conducting pins (note: pins extend out of page toward reader).

Figure 34 is a photograph showing the copper electrode plate placed on top of an assembled Caco-2 membrane bottom well and receiver tray.

Figure 35 is a photograph showing the assembled embodiment of Figure 34, i.e., electrode head, copper electrode plate with pins, Caco-2 membrane bottom well, and Caco-2 receiver tray.

Figure 36 depicts a novel electrode embodiment that comprises a dielectric disc sandwiched between two conductive discs. Figure 36A shows an expanded view of the novel electrode embodiment. Figure 36B shows the novel electrode embodiment electrically connected to a concentric lead. Figure 36C shows the novel electrode embodiment electrically connected to edge leads.

DETAILED DESCRIPTION OF THE INVENTION

The present invention provides equipment and techniques to implement electric field stimulation (EFS) of cells while monitoring a biological response of the cells. Preferably, the biological response is monitored by fluorescence detection. The cells are grown and/or attached to specially designed substrates such as, *e.g.*, glass slides which contain preferably transparent, electrically conductive electrodes or multiwell tissue culture plates containing electrodes so disposed that when a preselected voltage is applied across the electrodes the transmembrane potential of cells within the wells of the multiwell tissue culture plates is altered.

In general terms, the present invention involves providing a substrate upon which living eukaryotic cells, preferably mammalian cells, are present where the cells express voltage-gated ion channels in their plasma membranes. Positive and negative electrodes are positioned either on or near the substrate so that when a voltage is applied through the electrodes the voltage-gated ion channels either open or close, thereby modulating the flow of at least one type of ion through the plasma membranes of the cells. This modulation of ion flow, or a change in membrane potential that results from the modulation of ion flow, is detected, either directly or indirectly, preferably by the use of fluorescent indicator compounds in the cells.

Collections of substances, *e.g.*, combinatorial libraries of small organic molecules, natural products, phage display peptide libraries, etc., are brought into contact with the voltage-gated ion channels in the plasma membranes of the cells and those substances that are able to affect the modulation of ion flow are identified. In this way, the present invention provides methods of screening for activators and inhibitors of voltage-gated ion channels. Such activators and inhibitors are expected to be useful as pharmaceuticals or as lead compounds from which pharmaceuticals can be developed by the usual processes of drug development, *e.g.*, medicinal chemistry.

During an applied extracellular electrical field, the cell membrane electrical capacitance will charge or discharge depending upon the polarity and orientation of the cell relative to the field. This results in a transient change in the transmembrane potential in a given patch of membrane. These transient changes in transmembrane potential will vary continuously around each cell depending upon the orientation of each patch of membrane relative to the applied field and the existing transmembrane potential. In each membrane patch, membrane potential will be perturbed away from the resting value by the applied external field. This change in membrane potential will in turn affect the proportion of open and closed voltage-gated ion channels in each local patch of membrane, which will affect the conductance of the voltage-gated ion channels and thus change the membrane potential further. This process is expected to vary around each cell such that, in any given cell, different patches of membrane and the embedded voltage-gated ion channels will experience different membrane potentials. In general, the membrane potential in a given patch of membrane will change at a rate that is proportional to its resistance ($1/\text{conductance}$) and its capacitance (C_m) such that $dV/dt = I/C_m$ where I is the total current flow ($I=V/R$) across the patch of membrane.

Figure 14 illustrates these concepts. For the sake of simplicity, the plasma membrane of the cell shown in Figure 14 is divided into four patches: left, top, right, and bottom. Current will flow between the electrodes if a voltage difference is applied. This will alter the cell membrane potential. If electrode 1 is positive and electrode 2 is negative, the membrane patch at the bottom of the cell will be hyperpolarized but the top patch will be depolarized. The left and right patches will "see" no change in membrane potential. If polarity is reversed, the opposite will occur.

In reality, of course, the cell's plasma membrane is a continuum of individual patches rather than the simplified system of four patches depicted in Figure 14. The applied voltage alters the membrane potentials of the various patches to many different values such that the embedded voltage-gated ion channels "sample" the many different potentials and are driven through their various conformational states. These include open states, closed states, high affinity drug bound states, and low affinity drug bound states.

Accordingly, the present invention provides a method for identifying modulators of the activity of a voltage-gated ion channel comprising:

10 (a) altering the transmembrane potential of at least a portion of the membrane of a cell expressing the voltage-gated ion channel by applying a voltage to the cells through extracellular electrodes while monitoring ion flow through the voltage-gated ion channel;

(b) exposing the cell in step (a) to a substance and monitoring ion flow through the voltage-gated ion channel;

15 (c) comparing the ion flow through the voltage-gated ion channel in step (a) and step (b);

where a difference in the ion flow through the voltage-gated ion channel in step (a) and step (b) indicates that the substance is a modulator of the voltage-gated ion channel.

A variation of the method comprises:

(a) dividing a plurality of cells expressing the voltage-gated ion channel into a control portion and a test portion;

25 (b) altering the transmembrane potential of the control portion of cells by applying a voltage to the cells through extracellular electrodes while monitoring ion flow through the voltage-gated ion channel;

(c) altering the transmembrane potential of the test portion of cells by applying the voltage to the cells through extracellular electrodes in the presence of a substance while monitoring ion flow through the voltage-gated ion channel;

30 (d) comparing the ion flow through the voltage-gated ion channel in step (b) and step (c);

where a difference in the ion flow through the voltage-gated ion channel in step (b) and step (c) indicates that the substance is a modulator of the voltage-gated ion channel.

For the sake of simplicity, the above methods are described in terms of “a” voltage-gated ion channel although those skilled in the art will understand that in actual practice the cells will express a plurality of the voltage-gated ion channels for which modulators are sought. Generally, each cell will express at least 10², 10³, 10⁴, 5 10⁵, 10⁶ or more molecules of the voltage-gated ion channel. Also, ion flow will be monitored through the plurality of the voltage-gated ion channels rather than through a single voltage-gated ion channel. Similarly, the methods will generally be practiced by employing a plurality of cells, even though the methods are described above in terms of “a” cell.

10 Generally, the methods of the present invention will be carried out on a substrate that is a modified version of a standard multiwell tissue culture plate or microtiter plate. Such substrates will have a place for the cells to be tested (generally the wells of the tissue culture plate or microtiter plate) and will have positive and negative electrodes (either built into the plate or nearby) in such an orientations with 15 respect to the cells that the electrodes can deliver a voltage potential that causes an alteration in the open/close state of the voltage-gated ion channels in the cells. The electrodes are extracellular, *i.e.*, they do not penetrate into or across the plasma membranes of the cells although they may touch the outside of the plasma membranes in certain embodiments. Extracellular electrodes do not include electrodes which 20 form a continuous connection with a cell’s interior, *e.g.*, patch/clamp electrodes.

Therefore, the present invention provides a method of identifying activators of a voltage-gated ion channel comprising:

- (a) providing a substrate upon which are living eukaryotic cells that express a plurality of the voltage-gated ion channels in their plasma membranes;
- 25 (b) providing positive and negative electrodes positioned either on or near the substrate such that when a preselected voltage is applied through the positive and negative electrodes the transmembrane electrical potential of the cells is altered such that at least a portion of the voltage-gated ion channels are closed;
- (c) applying the preselected voltage through the positive and 30 negative electrodes;
- (d) determining a control value for the flow of ions through the voltage-gated ion channels of the cells in step (c);
- (e) exposing the cells of step (c) to a substance for a period sufficient and under conditions such that a detectable number of the portion of the

voltage-gated ion channels that are closed become open and allow ion flow through the detectable number of voltage-gated ion channels if the substance is an activator of the voltage-gated ion channels;

- (f) determining a test value for the flow of ions through the voltage-gated ion channels of the cells of step (e);
 - (g) comparing the control value to the test value;
- where if the control value is less than the test value, then the substance is an activator of the voltage-gated ion channel.

The above-described method can be easily modified to provide a method for identifying inhibitors of the voltage-gated ion channel. The voltage applied through the electrodes is preselected such that it alters the electrical field around the cells and consequently alters the transmembrane electrical field. This in turn changes the states of the embedded voltage-gated ion channels such that instead of the voltage-gated ion channels being closed, the voltage-gated ion channels may open. Substances are then screened for the ability to close or inhibit the channels.

Accordingly, the present invention provides a method of identifying inhibitors of a voltage-gated ion channel comprising:

- (a) providing a substrate upon which are living eukaryotic cells that express a plurality of the voltage-gated ion channels in their plasma membranes;
- (b) providing positive and negative electrodes positioned either on or near the substrate such that when a preselected voltage is applied through the positive and negative electrodes the transmembrane electrical potential of the cells is altered such that at least a portion of the voltage-gated ion channels are open;
- (c) applying the preselected voltage through the positive and negative electrodes;
- (d) determining a control value for the flow of ions through the voltage-gated ion channels of the cells in step (c);
- (e) exposing the cells of step (c) to a substance for a period sufficient and under conditions such that a detectable number of the portion of the voltage-gated ion channels that are open become closed and restrict ion flow through the detectable number of voltage-gated ion channels if the substance is an inhibitor of the voltage-gated ion channels;
- (f) determining a test value for the flow of ions through the voltage-gated ion channels of the cells of step (e);

(g) comparing the control value to the test value;

where if the control value is greater than the test value, then the substance is an inhibitor of the voltage-gated ion channel.

In the above-described method for identifying activators, the terms “a
5 portion of the voltage-gated ion channels are closed” and “a detectable number” are related and have relative rather than absolute values. Similarly, in the above-described method for identifying inhibitors, the terms “a portion of the voltage-gated ion channels are open” and “a detectable number” are also related and have relative rather than absolute values. What is meant is that a portion of the voltage-gated ion
10 channels will be open or closed such that when the substance acts on the channels, a change in the open/closed state of a sufficient number of channels (*i.e.*, “a detectable number”) occurs such that a difference in ion flow that is large enough to be measured by the detection system employed takes place. There is no need to determine the actual number of ion channels that constitutes the “portion” of voltage-gated ion
15 channels that are closed or open or the “detectable number” so long as the difference in ion flow can be measured. The actual portion of channels that will be open or closed as well as the actual value of “detectable number” in order for the methods to be practiced will depend on such variables as the channel that is being studied, the concentrations of the substances tested, the nature of the detection system for ion
20 flow, and so forth. Adjusting the voltage applied through the electrodes to take into account such variables so that control and test values can be obtained is a matter of routine experimentation in which the skilled artisan will be guided by knowledge in the art such as, *e.g.*, the known voltage dependence of the open/close transition of the voltage-gated ion channel under study, the nature and sensitivity of the detection
25 system employed to monitor the flow of ions, the level of expression of the ion channel in the cells, and so forth.

The electrodes can be arranged in a variety of ways in order to provide for the proper stimulus. A number of arrangements are described herein and illustrated in the accompanying figures. These include arrangements where the cells
30 are present in wells in the substrate and:

- (a) both a positive and negative electrode is present in each well;
- (b) one electrode is present in the well and the other electrode enters the fluid medium in the well from above without touching the sides or bottom of the well;

- (c) the electrodes form part of the sides or bottom of the wells;
- (d) a pattern of interdigitating electrodes has been formed on the

surface of the substrate and at least some of the cells are positioned between the interdigitating branches of the positive and negative electrodes.

5 The skilled person will recognize that it is generally beneficial to run controls together with the methods described herein. For example, it will usually be helpful to have a control in which the substances are tested in the methods against cells that preferably are essentially identical to the cells that are used in the methods except that these cells would not express the voltage-gated ion channels of interest. In
10 this way it can be determined that substances which are identified by the methods are really exerting their effects through the voltage-gated ion channels of interest rather than through some unexpected non-specific mechanism. One possibility for such control cells would be to use non-recombinant parent cells where the cells of the actual experiment express the voltage-gated ion channels of interest due to the
15 recombinant expression of those voltage-gated ion channels of interest.

Other types of controls would involve taking substances that are identified by the methods of the present invention as activators or inhibitors of voltage-gated ion channels of interest and testing those substances in the methods of the prior art in order to confirm that those substances are also activators and inhibitors
20 when tested in those prior art methods.

One skilled in the art would recognize that, where the present invention involves comparing control values for the flow of ions to test values for the flow of ions and determining whether the control values are greater or less than the test values, a non-trivial difference is sought. For example, if in the methods of
25 identifying inhibitors, the control value were found to be 1% greater than the test value, this would not indicate that the substance is an inhibitor. Rather, one skilled in the art would attribute such a small difference to normal experimental variance. What is looked for is a significant difference between control and test values. For the purposes of this invention, a significant difference fulfills the usual requirements for a
30 statistically valid measurement of a biological signal. For example, depending upon the details of the experimental arrangement, a significant difference might be a difference of at least 10%, preferably at least 20%, more preferably at least 50%, and most preferably at least 100%.

One skilled in the art would understand that the cells that give rise to the control values need not be physically the same cells that give rise to the test values, although that is possible. What is necessary is that the cells that give rise to the control values be substantially the same type of cell as the cells that give rise to the test values. A cell line that has been transfected with and expresses a certain voltage-gated ion channel could be used for both the control and test cells. Large numbers of such cells could be grown and a portion of those cells could be exposed to the substance and thus serve as the cells giving rise to the test value for ion flow while a portion would not be exposed to the substance and would thus serve as the cells giving rise to the control value for ion flow. No individual cell itself would be both control and test cell but the virtual identity of all the cells in the cell line ensures that the methods would nevertheless be reliable.

Accordingly, the present invention provides a method of identifying activators of a voltage-gated ion channel comprising:

- (a) providing a substrate upon which are living eukaryotic cells that express a plurality of the voltage-gated ion channels in their plasma membranes;
- (b) providing positive and negative electrodes positioned either on or near the substrate such that when a preselected voltage is applied through the positive and negative electrodes the transmembrane electrical potential of the cells is altered such that at least a portion of the voltage-gated ion channels are closed;
- (c) applying the preselected voltage through the positive and negative electrodes to a control sample of the cells;
- (d) determining a control value for the flow of ions through the voltage-gated ion channels of the control sample of the cells in step (c);
- (e) applying the preselected voltage through the positive and negative electrodes to a test sample of the cells while exposing the test sample of the cells to a substance for a period sufficient and under conditions such that a detectable number of the portion of the voltage-gated ion channels that are closed in the test sample become open and allow ion flow through the detectable number of voltage-gated ion channels if the substance is an activator of the voltage-gated ion channels;
- (f) determining a test value for the flow of ions through the voltage-gated ion channels of the test sample of cells of step (e);
- (g) comparing the control value to the test value;

where if the control value is less than the test value, then the substance is an activator of the voltage-gated ion channel.

Similarly, the present invention provides a method of identifying inhibitors of a voltage-gated ion channel comprising:

- 5 (a) providing a substrate upon which are living eukaryotic cells that express a plurality of the voltage-gated ion channels in their plasma membranes;
- (b) providing positive and negative electrodes positioned either on or near the substrate such that when a preselected voltage is applied through the positive and negative electrodes the transmembrane electrical potential of the cells is
10 altered such that at least a portion of the voltage-gated ion channels are open;
- (c) applying the preselected voltage through the positive and negative electrodes to a control sample of the cells;
- (d) determining a control value for the flow of ions through the voltage-gated ion channels of the control sample of the cells in step (c);
- 15 (e) applying the preselected voltage through the positive and negative electrodes to a test sample of the cells while exposing the test sample of the cells to a substance for a period sufficient and under conditions such that a detectable number of the portion of the voltage-gated ion channels that are open in the test sample become closed and restrict ion flow through the detectable number of voltage-
20 gated ion channels if the substance is an inhibitor of the voltage-gated ion channels;
- (f) determining a test value for the flow of ions through the voltage-gated ion channels of the test sample of cells of step (e);
- (g) comparing the control value to the test value;
- where if the control value is greater than the test value, then the
25 substance is an inhibitor of the voltage-gated ion channel.

“Substances” can be any substances that are generally screened in the pharmaceutical industry during the drug development process. For example, substances may be low molecular weight organic compounds (*e.g.*, having a molecular weight of less than about 1,000 daltons); RNA, DNA, antibodies, peptides,
30 or proteins.

The conditions under which cells are exposed to substances in the methods described herein are conditions that are typically used in the art for the study of protein-ligand interactions: *e.g.*, physiological pH; salt conditions such as those represented by such commonly used buffers as PBS or in tissue culture media; a

temperature of about 4°C to about 55°C; incubation times of from several seconds to several hours. Generally, the cells are present in wells in the substrate and the substances are added directly to the wells, optionally after first washing away the media in the wells.

5 Determining the values of ion flow in the methods of the present invention can be accomplished through the use of fluorescent indicator compounds. One type of fluorescent indicator compound is sensitive to the level of intracellular calcium ions in the cells used in the present invention. This type of fluorescent indicator compound can be used when the methods are directed to those voltage-gated
10 ion channels whose activity results in a change in intracellular calcium levels. Such voltage-gated ion channels include not only voltage-gated calcium channels but also other types of voltage-gated ion channels where the activity of those channels is naturally or can be coupled to changes in intracellular calcium levels. Many types of voltage-gated potassium channels can be so coupled. When using this approach to
15 study a voltage-gated ion channel of interest that is not a voltage-gated calcium channel, it may be desirable to engineer the cells employed so as to recombinantly express voltage-gated calcium channels that are coupled to the voltage-gated ion channel of interest.

Fluorescent indicator compounds suitable for measuring intracellular
20 calcium levels include various calcium indicator dyes (*e.g.*, fura-2, fluo-3, indo-1, Calcium Green; see Velicelebi et al., 1999, *Meth. Enzymol.* 294:20-47).

Calcium indicator dyes are substances which show a change in a fluorescent characteristic upon binding calcium, *e.g.*, greatly increased intensity of fluorescence and/or a change in fluorescent spectra (*i.e.*, a change in emission or
25 excitation maxima). Fluo-3, fura-2, and indo-1 are commonly used calcium indicator dyes that were designed as structural analogs of the highly selective calcium chelators ethylene glycol-bis(β -aminoethyl ether) N,N,N',N'-tetraacetic acid (EGTA) and 1,2-bis(2-aminophenoxy) ethane-N,N,N',N'-tetraacetic acid (BAPTA). The fluorescence intensity from fluo-3 increases by more than 100-fold upon binding of calcium.
30 While the unbound dye exhibits very little fluorescence, calcium-bound fluo-3 shows strong fluorescence emission at 526 nm. Fura-2 is an example of a dye that exhibits a change in its fluorescence spectrum upon calcium binding. In the unbound state, fura-2 has an excitation maximum of 362 nm. This excitation maximum shifts to 335 nm upon calcium binding, although there is no change in emission maximum. Binding of

calcium to fura-2 can be monitored by excitation at the two excitation maxima and determining the ratio of the amount of fluorescence emission following excitation at 362 nm compared to the amount of fluorescence emission following excitation at 335 nm. A smaller ratio (*i.e.*, less emission following excitation at 362 nm) indicates that
5 more fura-2 is bound to calcium, and thus a higher internal calcium concentration in the cell.

The use of calcium indicator dyes entails loading cells with the dye, a process which can be accomplished by exposing cells to the membrane-permeable acetoxymethyl esters of the dyes. Once inside the plasma membrane of the cells,
10 intracellular esterases cleave the esters, exposing negative charges in the free dyes. This prevents the free dyes from crossing the plasma membrane and thus leaves the free dyes trapped in the cells. Measurements of fluorescence from the dyes are then made, the cells are treated in such a way that the internal calcium concentration is changed (*e.g.*, by exposing cells to an activator or inhibitor of a voltage-gated ion
15 channel), and fluorescence measurements are again taken.

Fluorescence from the indicator dyes can be measured with a luminometer or a fluorescence imager. One preferred detection instrument is the Fluorometric Imaging Plate Reader (FLIPR) (Molecular Devices, Sunnyvale, CA). The FLIPR is well suited to high throughput screening using the methods of the
20 present invention as it incorporates integrated liquid handling capable of simultaneously pipetting to 96 or 384 wells of a microtiter plate and rapid kinetic detection using a argon laser coupled to a charge-coupled device imaging camera.

A typical protocol for use of calcium indicator dyes would entail plating cells expressing a voltage-gated ion channel of interest into an appropriate
25 substrate (*e.g.*, clear, flat-bottom, black-wall 96 well plates that have a suitable arrangement of positive and negative electrodes) and allowing the cells to grow overnight in standard tissue culture conditions (*e.g.*, 5% CO₂, 37°C). The cells are generally plated at a density of about 10,000 to 100,000 cells per well in appropriate growth medium. On the day of the assay, growth medium is removed and dye
30 loading medium is added to the wells.

If the calcium indicator dye is fluo-3, *e.g.*, dye loading medium could be prepared by solubilizing 50 µg of fluo-3-AM ester (Molecular Probes F-1242) in 22 µl DMSO to give a 2 mM dye stock. Immediately before loading the cells, 22 µl 20% pluronic acid (Molecular Probes P-3000) is added to the dye. The tube

containing the dye is mixed with a vortex mixer and 42 ml of the dye/pluronic acid solution is added to 10.5 ml of Hanks Balanced Salt Solution (Gibco/BRL Cat # 14025-076) with 20 mM HEPES (Gibco/BRL Cat # 1560-080), 2.5 mM probenecid (Sigma Cat # P-8761), and 1% fetal bovine serum (Gibco/BRL Cat # 26140-087; not BSA)). The dye and the loading medium are mixed by repeated inversion (final dye concentration about 4 μ M).

Growth medium can be removed from the cells by washing (wash medium is Hanks Balanced Salt Solution (Gibco/BRL Cat # 14025-076) with 20 mM HEPES (Gibco/BRL Cat # 1560-080), 2.5 mM probenecid (Sigma Cat # P-8761), and 0.1% bovine serum albumin (Sigma Cat # A-9647; not FBS) three times, leaving 100 μ l residual medium in the wells after the fourth wash. Then 100 μ l of the dye in the loading medium is added to each well. The cells are then incubated for 60 minutes to allow for dye loading.

Following dye loading, fluorescent measurements of the cells are taken prior to exposure of the cells to substances that are to be tested. The cells are then exposed to the substances and those substances that cause a change in a fluorescent characteristic of the dye are identified. The measuring instrument can be a fluorescent plate reader such as the FLIPR (Molecular Devices). Substances that cause a change in a fluorescent characteristic in the test cells but not the control cells are possible activators or inhibitors of the voltage-gated ion channel.

The exact details of the procedure outlined above are meant to be illustrative. One skilled in the art would be able to optimize experimental parameters (cell number, dye concentration, dye loading time, temperature of incubations, cell washing conditions, and instrument settings, etc.) by routine experimentation depending on the particular relevant experimental variables (e.g., type of cell used, identity of dye used). Several examples of experimental protocols that can be used are described in Veliçelebi et al., 1999, Meth. Enzymol. 294:20-47. Other suitable instrumentation and methods for measuring transmembrane potential changes via optical methods includes microscopes, multiwell plate readers and other instrumentation that is capable of rapid, sensitive ratiometric fluorescence detection. For example, the VIPR (Aurora Biosciences, San Diego, CA) is an integrated liquid handler and kinetic fluorescence reader for 96-well and greater multiwell plates. The VIPR reader integrates an eight channel liquid handler, a multiwell positioning stage and a fiber-optic illumination and detection system. The system is designed to measure fluorescence from a column of eight wells simultaneously before, during and after the introduction of liquid

sample obtained from another microtiter plate or trough. The VIPR reader excites and detects emission signals from the bottom of a multiwell plate by employing eight trifurcated optical bundles (one bundle for each well). One leg of the trifurcated fiber is used as an excitation source, the other two legs of the trifurcated fiber being used to
5 detect fluorescence emission. A ball lens on the end of the fiber increases the efficiency of light excitation and collection. The bifurcated emission fibers allow the reader to detect two emission signals simultaneously and are compatible with rapid signals generated by the FRET-based voltage dyes.

Photomultiplier tubes then detect emission fluorescence, enabling sub-second
10 emission ratio detection.

In particular embodiments, the calcium indicator dye is selected from the group consisting of: fluo-3, fura-2, fluo-4, fluo-5, calcium green-1, Oregon green, 488 BAPTA, SNARF-1, and indo-1.

15 In particular embodiments, the change in fluorescent characteristic is an increase in intensity of a fluorescence emission maximum. In other embodiments, the change in fluorescent characteristic is a shift in the wavelength of an absorption maximum.

In particular embodiments, the cells naturally express the voltage-gated
20 ion channel of interest and/or calcium channels. In other embodiments, the cells do not naturally express the voltage-gated ion channel of interest and/or calcium channels but instead have been transfected with expression vectors that encode the voltage-gated ion channel of interest and/or calcium channels so that the cells recombinantly express the voltage-gated ion channel of interest and/or calcium
25 channels. Transfection is meant to include any method known in the art for introducing expression vectors into the cells. For example, transfection includes calcium phosphate or calcium chloride mediated transfection, lipofection, infection with a retroviral construct, and electroporation.

An alternative to the use of calcium indicator dyes is the use of the
30 aequorin system. The aequorin system makes use of the protein apoequorin, which binds to the lipophilic chromophore coelenterazine forming a combination of apoequorin and coelenterazine that is known as aequorin. Apoequorin has three calcium binding sites and, upon calcium binding, the apoequorin portion of aequorin

changes its conformation. This change in conformation causes coelenterazine to be oxidized into coelenteramide, CO₂, and a photon of blue light (466 nm). This photon can be detected with suitable instrumentation.

Since the gene encoding apoaequorin has been cloned (U.S. Patent No. 5,541,309; U.S. Patent No. 5,422,266; U.S. Patent No. 5,744,579; Inouye et al., 1985, Proc. Natl. Acad. Sci. USA 82:3154-3158; Prasher et al., 1985, Biochem. Biophys. Res. Comm. 126:1259-1268), apoaequorin can be recombinantly expressed in cells in which it is desired to measure the intracellular calcium concentration. Alternatively, existing cells that stably express recombinant apoaequorin can be used. Such cells derived from HEK293 cells and CHO-K1 cells are described in Button & Brownstein, 1993, Cell Calcium 14:663-671. For example, the HEK293/aeq17 cell line can be used as follows.

The HEK293/aeq17 cells are grown in Dulbecco's Modified Medium (DMEM, GIBCO-BRL, Gaithersburg, MD, USA) with 10% fetal bovine serum (heat inactivated), 1 mM sodium pyruvate, 500 µg/ml Geneticin, 100 µg/ml streptomycin, 100 units/ml penicillin. Expression vectors encoding the voltage-gated ion channel of interest as well as, optionally, the desired voltage-gated calcium channel subunits (α 1A, α 1B, α 1C, α 1D, α 1E, α 1G, α 1H, α 1I, α 2 δ , β 1, β 2, β 3, β 4, etc.) can be transfected into the HEK293/aeq17 cells by standard methods in order to express the desired voltage-gated ion channel subunits and voltage-gated calcium channel subunits in the HEK293/aeq17 cells. The cells are washed once with DMEM plus 0.1 % fetal bovine serum, and then charged for one hour at 37°C /5% CO₂ in DMEM containing 8 µM coelenterazine cp (Molecular Probes, Eugene, OR, USA) and 30 µM glutathione. The cells are then washed once with Versene (GIBCO-BRL, Gaithersburg, MD, USA), detached using Enzyme-free cell dissociation buffer (GIBCO-BRL, Gaithersburg, MD, USA), diluted into ECB (Ham's F12 nutrient mixture (GIBCO-BRL) with 0.3 mM CaCl₂, 25 mM HEPES, pH7.3, 0.1% fetal bovine serum). The cell suspension is centrifuged at 500 x g for 5 min. The supernatant is removed, and the pellet is resuspended in 10 ml ECB. The cell density is determined by counting with a hemacytometer and adjusted to 500,000 cells/ml in ECB. The substances to be tested are diluted to the desired concentrations in ECB and aliquoted into the assay plates, preferably in triplicate, at 0.1 ml/well. The cell suspension is injected at 0.1 ml/well, read and integrated for a total of 400 readings using a luminometer (Luminoskan Ascent, Labsystems Oy, Helsinki, Finland).

Alternatively, the cells may first be placed into the assay plates and then the substances added. Data are analyzed using the software GraphPad Prism Version 3.0 (GraphPad Software, Inc., San Diego, CA, USA).

It will be understood by those skilled in the art that the procedure
5 outlined above is a general guide in which the various steps and variables can be modified somewhat to take into account the specific details of the particular assay that is desired to be run. For example, one could use semisynthetic coelenterazine (Shimomura, 1989, *Biochem. J.* 261:913-920; Shimomura et al., 1993, *Cell Calcium* 14:373-378); the time of incubation of the cells with coelenterazine can be varied
10 somewhat; somewhat greater or lesser numbers of cells per well can be used; and so forth.

For reviews on the use of aequorin, see Créton et al., 1999, *Microscopy Research and Technique* 46:390-397; Brini et al., 1995, *J. Biol. Chem.* 270:9896-9903; Knight & Knight, 1995, *Meth. Cell. Biol.* 49:201-216. Also of interest may be
15 U.S. Patent No. 5,714,666 which describes methods of measuring intracellular calcium in mammalian cells by the addition of coelenterazine co-factors to mammalian cells that express apoequorin.

Another way to measure ion flow is to monitor changes in transcription that result from the activity of voltage-gated ion channels by the use of transcription
20 based assays. Transcription-based assays involve the use of a reporter gene whose transcription is driven by an inducible promoter whose activity is regulated by a particular intracellular event such as, *e.g.*, changes in intracellular calcium levels, that are caused by the activity of a voltage-gated ion channel. Transcription-based assays are reviewed in Rutter et al., 1998, *Chemistry & Biology* 5:R285-R290.
25 Transcription-based assays of the present invention rely on the expression of reporter genes whose transcription is activated or repressed as a result of intracellular events that are caused by the interaction of a activator or inhibitor with a voltage-gated ion channel.

An extremely sensitive transcription-based assay is disclosed in
30 Zlokarnik et al., 1998, *Science* 279:84-88 (Zlokarnik) and also in U.S. Patent No. 5,741,657. The assay disclosed in Zlokarnik and U.S. Patent No. 5,741,657 employs a plasmid encoding β -lactamase under the control of an inducible promoter. This plasmid is transfected into cells together with a plasmid encoding a receptor for which it is desired to identify agonists. The inducible promoter on the β -lactamase is chosen

so that it responds to at least one intracellular signal that is generated when an agonist binds to the receptor. Thus, following such binding of agonist to receptor, the level of β -lactamase in the transfected cells increases. This increase in β -lactamase is measured by treating the cells with a cell-permeable dye that is a substrate for cleavage by β -lactamase. The dye contains two fluorescent moieties. In the intact dye, the two fluorescent moieties are physically linked, and thus close enough to one another that fluorescence resonance energy transfer (FRET) can take place between them. Following cleavage of the dye into two parts by β -lactamase, the two fluorescent moieties are located on different parts, and thus can diffuse apart. This increases the distance between the fluorescent moieties, thus decreasing the amount of FRET that can occur between them. It is this decrease in FRET that is measured in the assay.

The assay described in Zlokarnik and U.S. Patent No. 5,741,657 can be modified for use in the methods of the present invention by using an inducible promoter to drive β -lactamase where the promoter is activated by an intracellular signal generated by the opening or closing of a voltage-gated ion channel. Cells expressing a voltage-gated ion channel and the inducible promoter-driven β -lactamase are placed in the apparatus of the present invention, where the open or closed state of the voltage-gated ion channels can be controlled. The cells are exposed to the cell-permeable dye and then exposed to substances suspected of being activators or inhibitors of the voltage-gated ion channel. Those substances that cause a change in the open or closed state of the voltage-gated ion channel are identified by their effect on the inducible promoter-driven β -lactamase and thus on FRET. The inducible promoter-driven β -lactamase is engineered with a suitable promoter so that β -lactamase is induced when the substance is either an activator or an inhibitor, depending upon the nature of the assay.

The flow of ions through voltage-gated ion channels can also be measured by measuring changes in membrane potential via the use of fluorescent voltage sensitive dyes. The changes in membrane potential will depend on the ion channels in the cell membrane. The resultant membrane potential will depend on the net properties of all the channels and the change caused by inhibiting (through a substance that is an inhibitor or antagonist) or activating (through a substance that is an activator or an agonist) the voltage-gated ion channel of interest. One knowledgeable in cellular and membrane biophysics and electrophysiology will

understand the directions of the changes in membrane potential since those changes depend on the ion channels present and the inhibition or activation of those channels by test substances. In many cases when using fluorescent voltage sensitive dyes, the experimental system can be calibrated by using known activators or inhibitors of the voltage-gated ion channel of interest.

The present invention therefore includes assays that monitor changes in ion flow caused by activators or inhibitors of voltage-gated ion channels based upon FRET between a first and a second fluorescent dye where the first dye is bound to one side of the plasma membrane of a cell expressing a voltage-gated ion channel of interest and the second dye is free to move from one face of the membrane to the other face in response to changes in membrane potential. In certain embodiments, the first dye is impenetrable to the plasma membrane of the cells and is bound predominately to the extracellular surface of the plasma membrane. The second dye is trapped within the plasma membrane but is free to diffuse within the membrane. At normal (*i.e.*, negative) resting potentials of the membrane, the second dye is bound predominately to the inner surface of the extracellular face of the plasma membrane, thus placing the second dye in close proximity to the first dye. This close proximity allows for the generation of a large amount of FRET between the two dyes. Following membrane depolarization, the second dye moves from the extracellular face of the membrane to the intracellular face, thus increasing the distance between the dyes. This increased distance results in a decrease in FRET, with a corresponding increase in fluorescent emission derived from the first dye and a corresponding decrease in the fluorescent emission from the second dye. See figure 1 of González & Tsien, 1997, *Chemistry & Biology* 4:269-277. See also González & Tsien, 1995, *Biophys. J.* 69:1272-1280 and U.S. Patent No. 5,661,035.

In certain embodiments, the first dye is a fluorescent lectin or a fluorescent phospholipid that acts as the fluorescent donor. Examples of such a first dye are: a coumarin-labeled phosphatidylethanolamine (*e.g.*, N-(6-chloro-7-hydroxy-2-oxo-2H--1-benzopyran-3-carboxamidoacetyl)-dimyristoylphosphatidylethanolamine) or N-(7-nitrobenz-2-oxa-1,3-diazol-4-yl)-dipalmitoylphosphatidylethanolamine); a fluorescently-labeled lectin (*e.g.*, fluorescein-labeled wheat germ agglutinin). In certain embodiments, the second dye is an oxonol that acts as the fluorescent acceptor. Examples of such a second dye are: bis(1,3-dialkyl-2-thiobarbiturate)trimethineoxonols (*e.g.*, bis(1,3-dihexyl-2-

thiobarbiturate)trimethineoxonol) or pentamethineoxonol analogues (*e.g.*, bis(1,3-dihexyl-2-thiobarbiturate)pentamethineoxonol; or bis(1,3-dibutyl-2-thiobarbiturate)pentamethineoxonol). See González & Tsien, 1997, Chemistry & Biology 4:269-277 for methods of synthesizing various dyes suitable for use in the present invention. In certain embodiments, the assay may comprise a natural carotenoid, *e.g.*, astaxanthin, in order to reduce photodynamic damage due to singlet oxygen.

The use of such fluorescent dyes capable of moving from one face of the plasma membrane to the other is especially appropriate when the methods of the present invention are directed to inwardly rectifying potassium channels. Activation of inwardly rectifying potassium channels results in increased potassium current flow across the plasma membrane. This increased current flow results in a hyperpolarization of the cell membrane that can be detected by use of the technique described above since such hyperpolarization will result in greater FRET.

A large number of possible combinations of types of substrates and electrodes; physical arrangement of electrodes; number, shape, and arrangement of wells for holding the cells are suitable for use in the present invention.

Figure 1 illustrates an embodiment of the invention where the electrodes are generally parallel wires or strips of conductive material such as gold. The electrodes lie on the surface of a glass substrate and, together with the spacers, form the walls of the wells. For clarity, only a single series of wells is shown in Figure 1. Generally, substantially the entire surface of the glass substrate would be covered by wells formed in the manner shown. Cells are placed in the wells and grown in suitable media until an appropriate number of cells is present in the wells. Alternatively, an appropriate number of cells may be placed into the wells and used without further growth.

Figure 2B illustrates an embodiment of the invention where the wells are cavities or depressions in the surface of the substrate, as in typical multiwell tissue culture plates. Each well has an electrode at the bottom of the well and another electrode that is aligned along a side of the well. The cells are shown in Figure 2B as attached at the bottom of the well but in certain embodiments the cells may be suspension cells dispersed in the fluid in the well.

Figure 2C illustrates an embodiment of the invention similar to that shown in Figure 2B except that in Figure 2C both electrodes are at the bottom of the wells.

Figure 3 illustrates an embodiment of the invention where an array of
5 interdigitating transparent electrodes has been chemically etched onto the surface of a glass substrate. The electrode array, comprising a comb of positive and negative electrodes, has been chemically etched onto an indium tin oxide (ITO) coated glass plate. The thin layer of ITO (about 200 Å to 2,000 Å, or 500 Å to 1,500 Å, preferably 1,200 Å thick) forms a transparent conductive coating on the surface of the glass.
10 Although not essential, it is preferred that the layer of ITO be thin enough to be transparent. The chemical etching process removes the ITO from selected areas, resulting in an array of transparent ITO electrodes bonded to the glass. Multiple reaction wells may be contained on a single glass plate by forming fluid retention wells at the different electrode array sites. The wells can be formed by attaching (*e.g.*,
15 gluing) a well frame to the glass substrate or by forming virtual wells on the glass plate by a method such as screening hydrophobic ink onto the plate.

Figure 4A and 4B illustrates an embodiment in which wells are formed by attaching a well frame onto the substrate.

Figure 6 illustrates an embodiment in which a droplet of fluid
20 containing cells that express a voltage-gated ion channel is sandwiched between two plates. The plates, which can be glass plates, are each coated with a thin layer of conductive material such as indium tin oxide (ITO). The layers of conductive material are connected to a pulse generator such that one layer functions as a positive electrode and the other layer functions as a negative electrode.

25 Figures 7 and 8 illustrate embodiments in which one of the electrodes enters the well from above. In Figures 9 and 10, both electrodes enter from above.

The substrates for use in the present invention may contain virtual wells. Virtual wells are formed when a surface is patterned to have relatively hydrophilic domains within relatively hydrophobic fields so that an aqueous sample is
30 physically constrained by surface tension to the more hydrophilic domains by the edges of the more hydrophobic fields. The hydrophilic domains can be small circles that form a pattern similar to the wells of a conventional microtiter plate. Virtual wells provide a location in which samples can be confined without the deep indentations found in conventional microtiter plates. Figure 5 illustrates a surface for

use in the present invention that is a derivatized glass surface upon which virtual wells have been formed and upon which a pattern of interdigitated electrodes has also been formed. Figure 3 shows an individual well from this surface. International Patent Publication WO 99/39829 describes virtual wells and how they can be made.

5 “Interdigitating” refers to an arrangement of positive and negative electrodes where the positive and negative electrodes contain branches that are arranged such that, if a line were drawn from one branch of a positive electrode to the adjacent branch of the positive electrode, the line would cross a branch of the negative electrode. Similarly, if a line were drawn from one branch of a negative electrode to
10 the adjacent branch of the negative electrode, the line would cross a branch of the positive electrode. Generally, each interdigitating positive or negative electrode has at least 2, or at least 4, or at least 10, or at least 20 interdigitating branches. An example of interdigitating electrodes is shown in Figure 3.

 Various additional arrangements of electrodes formed from conductive
15 materials on glass substrates are possible. One arrangement has the positive and negative electrodes formed on two parallel glass substrates. For example, instead of having the positive and negative electrodes on a single glass substrate, two ITO coated glass substrates can be utilized by placing the glass substrates parallel to one another and placing the biologic fluid containing the cells in the gap between the glass
20 substrates. In this arrangement, one conductive glass substrate serves as the positive electrode while the second glass substrate serves as the negative electrode. The electrode field is formed at a right angle to the surface of the plates. This arrangement would allow fluid containing the cells to be either dispensed in between the plates or drawn into the gap via capillary action. The detector’s light beam would enter
25 perpendicular to the glass substrates and pass into the gap between the glass substrates, illuminating the fluid and cells. The fluorescence transmission from the cells would be collected by the detector in a similar manner. Figure 6 illustrates one version of this arrangement. Another version is shown in Figure 13 where an embodiment comprising two ITO-coated plates each containing multiple virtual wells is depicted. The ITO forms the bottom of the wells as well as the electrodes.
30

 Another arrangement has the positive and negative electrodes formed by a single glass substrate and a reference electrode. This arrangement utilizes a single glass substrate coated with a conductive material such as ITO as one electrode. A well holding the biological fluid and cells is formed on the surface of the

conductive material coating the glass substrate. A wire or similar conducting member placed into the well serves as the second electrode. Figure 7 illustrates a single well of a version of this arrangement. Figure 12 depicts this type of arrangement as it is usually practiced, in a multiwell format. Figure 15 shows a modification of this arrangement where one electrode is a highly conductive metal grid that is in contact with the ITO layer.

Another arrangement has the single conductive glass substrate acting as the conductor to the current generated by a positive and negative electrode pair placed in adjacent wells. See Figure 16A-D. This arrangement does not use a grounding grid. The current flows from a first electrode in a first well through the ITO bottom of the first well to the ITO bottom of an adjacent second well and through a second electrode in the second well. Adjacent electrodes are alternately positive and negative. See Figure 16A and 16C.

In certain embodiments using interdigitating electrodes, the spacing and width of the branches of the electrodes are on the same order of magnitude as the size of individual cells. Cells may be grown and attached to the substrate in such a manner that, if a cell attaches between a pair of positive and negative electrode branches, a lower applied stimulus pulse can be utilized. The advantage of this close electrode spacing is that it results in less shunting of the stimulus current pulse through the fluid medium and less fluid heating while stimulating the cells. The use of transparent interdigitating electrodes offers the advantage of passing light from a fluorescent emission light source through the preferably glass substrate and transparent electrodes onto the cell and light passage of the fluorescence signal back to the light detector. While making the electrodes from a transparent material such as indium tin oxide (ITO) has advantages in certain embodiments, the electrodes may also be made from non-transparent conductive materials such as platinum, silver, or gold. If the electrode material is not transparent, fluorescence measurements are still possible because light can pass through the glass in between the electrodes.

Regardless of the arrangement of electrodes, stimulus pulses are generated by a pulse generator and applied to either a single well electrode array or to multiple well electrode arrays. Various commercial pulse generators can be utilized that permit waveform generation and amplitude adjustment. Constant voltage or constant current waveforms can be applied to the electrodes. Commercially available

power supplies that can be used in the present invention include the STG 1004 or STG 1008 Stimulus Generator or the National Instruments PCI 6713 8 channel pcb.

In using the pulse generator to stimulate the cells, particular attention should be paid to the amplitude, pulse width, and polarity used. For certain extreme field strengths, electroporation of the biological membrane can occur, and this should be avoided. When changing the external electrical field, the desired goal is a change in the trans-membrane field (V_m) by less than approximately ± 100 mV. As such the amount of charge added or removed from the cell membrane capacitance is critical. Adjustment of the pulse amplitude and duration is necessary to ensure a change in V_m without electroporation of the cells. Typically the voltage changes across the electrodes may be on the order of ± 10 volts, preferably less than ± 5 volts, and if possible less than ± 1 volt. These values can be adjusted empirically, by routine experimentation, in order to optimize the cellular membrane potential change without electroporation of the cell membrane. In general, the amount of charge change on the cell membrane will depend upon the local field changes, which depend upon the electrical current. Adjusting the area (the current-time integral) of the applied current as determined by the change in external electric field can be readily optimized empirically. In general, if the goal is to stimulate a cellular action potential, the pulse duration will be kept brief and the amplitude will be increased up to a point that exceeds the threshold for action potential generation. This will be affected by the relative levels of ion channels expressed in the cells and will vary accordingly, requiring empirical adjustment. A typical value might be a pulse duration of 1 millisecond and a pulse amplitude of 5 volts; this might be varied to increase the duration to 2 milliseconds and decrease the amplitude to 2.5 volts, or to decrease the duration and increase the amplitude, etc. In general, there is an inverse parabolic relationship between the duration and the amplitude of the applied pulse, where the area of the applied current-time integral remains constant. Because ion channel kinetics and action potentials can be rapid and brief, minimizing the pulse duration is useful. These parameters will also depend upon the manufactured electrodes, their capacitance and resistance, the geometrical relationship to the cells, the ionic strength and composition of the solutions used, and the electrical coupling to the cells. Because of these many variables, an empirical approach based upon the above guidelines is best.

Electrode arrangements can be adapted to 12-well, 24-well, 96-well, 384-well, 1,536-well, 3,456-well, and other plate formats, permitting the present invention to be used in high throughput screening applications.

In embodiments of the invention such as that illustrated in Figure 12
5 where multiple wells are present in the substrate and each well has an electrode associated with it, the stimulus delivered to each well through the electrodes can be individually controlled by the application of suitable software that governs the pulse generator. Such software is well known in the art or can be readily designed by one skilled in the art.

10 Particular embodiments of the present invention employ an arrangement of electrodes and wells on a substrate such that the substrate has the same form factor as a typical multiwell tissue culture plate that is used for high throughput screening, *e.g.*, a 96 well plate. The spacing of the wells on the substrate can be such that the center-to-center distances of the wells on the substrate is the same
15 as the typical center-to-center distances between wells on typical 96 well plates that are used for high throughput screening. This facilitates the use of the present invention with current equipment used in high throughput screening such as plate handlers, detectors, automatic pipettors, etc. Substrates can be manufactured by modifying the well-known manufacturing processes generally used to make multiwell
20 tissue culture plates by adding electrodes to the plates according to one of the configurations of electrodes disclosed herein.

In particular embodiments of the present invention, the substrate is not silicon or a field effect transistor.

In particular embodiments of the present invention, cells are utilized
25 that have been transfected with expression vectors comprising DNA that encodes a voltage-gated ion channel. Preferably, the cells do not naturally express corresponding voltage-gated ion channels. For example, if the expression vectors direct the expression of a voltage-gated calcium channel, the cells will not naturally express voltage-gated calcium channels. Alternatively, if the cells naturally express
30 corresponding voltage-gated ion channels, those corresponding voltage-gated ion channels can be distinguished from the transfected voltage-gated ion channels in some manner, *e.g.*, by the use of appropriate inhibitors, by manipulation of membrane potential. A preferred cell line for use in the present invention is the HEK293 cell line (ATCC 1573) since this cell line naturally expresses endogenous potassium

channels, which may be beneficial for electrical field stimulation experiments with channels that cause membrane potential depolarization (*e.g.*, sodium or calcium channels).

Cells are generally eukaryotic cells, preferably mammalian cells. The cells may be grown to the appropriate number on the substrates or they may be placed on the substrate and used without further growth. The cells may be attached to the substrate or, in those embodiments where the cells are placed or grown in wells, the cells may be suspension cells that are suspended in the fluid in the wells. Primary cells or established cell lines may be used.

Suitable cells for transfection with expression vectors that direct the expression of voltage-gated ion channels include but are not limited to cell lines of human, bovine, porcine, monkey and rodent origin. The cells may be adherent or non-adherent. Cells and cell lines which are suitable and which are widely available, include but are not limited to: L cells L-M(TK⁻) (ATCC CCL 1.3), L cells L-M (ATCC CCL 1.2), HEK293 (ATCC CRL 1573), Raji (ATCC CCL 86), CV-1 (ATCC CCL 70), COS-1 (ATCC CRL 1650), COS-7 (ATCC CRL 1651), CHO-K1 (ATCC CCL 61), 3T3 (ATCC CCL 92), NIH/3T3 (ATCC CRL 1658), HeLa (ATCC CCL 2), C127I (ATCC CRL 1616), BS-C-1 (ATCC CCL 26), MRC-5 (ATCC CCL 171), CPAE (ATCC CCL 209), Saos-2 (ATCC HTB-85), ARPE-19 human retinal pigment epithelium (ATCC CRL-2302), GH3 cells, and primary cardiac myocytes.

A variety of voltage-gated ion channels may be used in the present invention. For example, voltage-gated sodium channels, voltage-gated potassium channels, and voltage-gated calcium channels are suitable.

In certain embodiments of the present invention, the cells used do not naturally express the voltage-gated ion channel of interest. Instead, DNA encoding the voltage-gated ion channel is transfected into cells in order to express the voltage-gated ion channel in the plasma membrane of the cells. DNA encoding voltage-gated ion channels can be obtained by methods well known in the art. For example, a cDNA fragment encoding a voltage-gated ion channel can be isolated from a suitable cDNA library by using the polymerase chain reaction (PCR) employing suitable primer pairs. The cDNA fragment encoding the voltage-gated ion channel can then be cloned into a suitable expression vector. Primer pairs can be selected based upon the known DNA sequence of the voltage-gated ion channel it is desired to obtain.

Suitable cDNA libraries can be made from cellular or tissue sources known to contain mRNA encoding the voltage-gated ion channel.

One skilled in the art would know that for certain voltage-gated ion channels, it is desirable to transfect, and thereby express, more than one subunit in order to obtain a functional voltage-gated ion channel. For example, N-type calcium channels are composed of a multisubunit complex containing at least an $\alpha 1B$, an $\alpha 2\delta$, and a $\beta 1$ subunit. On the other hand, T-type calcium channels are functional with only a single subunit, *e.g.*, $\alpha 1G$, $\alpha 1H$, or $\alpha 1I$. Common knowledge in the art of the subunit composition of a voltage-gated ion channel of interest will lead the skilled artisan to express the correct subunits in the transfected cells.

One skilled in the art could use published voltage-gated ion channel sequences to design PCR primers and published studies of voltage-gated ion channel expression to select the appropriate sources from which to make cDNA libraries in order to obtain DNA encoding the voltage-gated ion channels. The following publications may be of use in this regard:

U.S. Patent No. 5,380,836 describes nucleic acid sequences encoding a rat cardiac voltage-gated sodium channel;

U.S. Patent No. 6,030,810 describes a number of voltage-gated, tetrodotoxin-sensitive sodium channels;

U.S. Patent No. 6,184,349 B1 discloses a human tetrodotoxin-resistant peripheral nerve voltage-gated sodium channel known as PN3; see also GenBank accession no. AF117907;

Isom et al., 1994, Neuron 12:1183-1194 discloses a rat voltage-gated sodium channel β subunit;

McClatchey et al., 1993, Hum. Molec. Gen. 2:745-749 discloses a human voltage-gated sodium channel $\beta 1$ subunit (hSCN $\beta 1$);

Isom et al., Science, 1992, 256:839-842 discloses a rat brain voltage-gated sodium channel $\beta 1$ subunit (rSCN $\beta 1$);

Misgeld et al., 1995, Prog. Neurobiol. 46:423-462; North, 1989, Br. J. Pharmacol. 98:13-23; Gahwiler et al., 1985, Proc. Natl. Acad. Sci USA 82:1558-1562; and Andrade et al., 1986, Science 234:1261-1265 disclose inwardly rectifying voltage-gated potassium channels that are suitable for use in the methods of the present invention.

U.S. Patent No. 5,874,236 and U.S. Patent No. 5,429,921 describe various $\alpha 1$ and β subunits of human voltage-gated calcium channels;

U.S. Patent No. 5,407,820 and U.S. Patent No. 5,710,250 describe $\alpha 2$ subunits of human voltage-gated calcium channels;

5 International Patent Publication WO 98/13490 describes a brain-specific P/Q-type human voltage-gated calcium channel involved in familial hemiplegic migraine;

Table 1 provides a list of ion channel genes that are suitable for use in the present invention.

10

TABLE 1

Some ion channel genes of interest for EFS experiments				
Symbol	Full Name	Cytogenetic Location	MIM Number	PubMed ID
SCN1	symbol withdrawn, see SCN1A			
SCN1A	sodium channel, voltage-gated, type I, alpha polypeptide	2q24	182389	8062593
SCN1B	sodium channel, voltage-gated, type I, beta polypeptide	19	600235	8394762
SCN2A1	sodium channel, voltage-gated, type II, alpha 1 polypeptide	2q22-q23	182390	1317301
SCN2A2	sodium channel, voltage-gated, type II, alpha 2 polypeptide	2q23-q24	601219	1317301
SCN2A	symbol withdrawn, see SCN2A1	-		
SCN2B	sodium channel, voltage-gated, type II, beta polypeptide	11q22-qter	601327	10198179
SCN3A	sodium channel, voltage-gated, type III, alpha polypeptide	2q24	182391	9589372
SCN4A	sodium channel, voltage-gated, type IV, alpha polypeptide	17q23-q25.3	603967	1654742
SCN4B	sodium channel, voltage-gated, type IV, beta polypeptide	reserved		
SCN5A	sodium channel, voltage-gated, type V, alpha polypeptide (long (electrocardiographic) QT syndrome 3)	3p21	600163	
SCN6A	sodium channel, voltage-gated, type VI, alpha polypeptide	2q21-q23	182392	10198179
SCN7A	symbol withdrawn, see SCN6A	-		
SCN8A	sodium channel, voltage gated, type VIII, alpha polypeptide	12q13.1	600702	7670495
SCN9A	sodium channel, voltage-gated, type IX, alpha polypeptide	2q24	603415	7720699

TABLE 1 (Continued)

Some ion channel genes of interest for EFS experiments				
Symbol	Full Name	Cytogenetic Location	MIM Number	PubMed ID
SCN10A	sodium channel, voltage-gated, type X, alpha polypeptide	3p21-p22	604427	9839820
SCN11A	sodium channel, voltage-gated, type XI, alpha polypeptide	3p21-p24	604385	10444332
SCN12A	sodium channel, voltage-gated, type XII, alpha polypeptide	3p23-p21.3		10623608
SCNN1	symbol withdrawn, see SCNN1A	-		
SCNN1A	sodium channel, nonvoltage-gated 1 alpha	12p13	600228	7896277
SCNN1B	sodium channel, nonvoltage-gated 1, beta (Liddle syndrome)	16p12.2-p12.1		600760
SCNN1D	sodium channel, nonvoltage-gated 1, delta	1p36.3-p36.2	601328	8661065
SCNN1G	sodium channel, nonvoltage-gated 1, gamma	16p12	600761	7490094
CACNA1A	calcium channel, voltage-dependent, P/Q type, alpha 1A subunit	19p13	601011	8825650
CACNA1B	calcium channel, voltage-dependent, L type, alpha 1B subunit	9q34	601012	8825650
CACNA1C	calcium channel, voltage-dependent, L type, alpha 1C subunit	12pter-p13.2	114205	1650913
CACNA1D	calcium channel, voltage-dependent, L type, alpha 1D subunit	3p14.3	114206	1664412
CACNA1E	calcium channel, voltage-dependent, alpha 1E subunit	1q25-q31	601013	8388125
CACNA1F	calcium channel, voltage-dependent, alpha 1F subunit	Xp11.23-p11.22	300110	9344658
CACNA1G	calcium channel, voltage-dependent, alpha 1G subunit	17q22	604065	9495342

TABLE 1 (Continued)

Some ion channel genes of interest for EFS experiments				
Symbol	Full Name	Cytogenetic Location	MIM Number	PubMed ID
CACNA1H	calcium channel, voltage-dependent, alpha 1H subunit	16p13.3		9670923
CACNA1I	calcium channel, voltage-dependent, alpha 1I subunit	22q12.3-13.2		10454147
CACNA1S	calcium channel, voltage-dependent, L type, alpha 1S subunit	1q31-q32	114208	7916735
CACNA2	symbol withdrawn, see CACNA2D1	-		
CACNA2D1	calcium channel, voltage-dependent, alpha 2/delta subunit 1	7q21-q22	114204	8188232
CACNA2D2	calcium channel, voltage-dependent, alpha 2/delta subunit 2	reserved		
CACNB1	calcium channel, voltage-dependent, beta 1 subunit	17q21-q22	114207	8381767
CACNB2	calcium channel, voltage-dependent, beta 2 subunit	10p12	600003	9254841
CACNB3	calcium channel, voltage-dependent, beta 3 subunit	12q13	601958	8119293
CACNB4	calcium channel, voltage-dependent, beta 4 subunit	2q22-q31	601949	9628818
CACNG1	calcium channel, voltage-dependent, gamma subunit 1	17q24	114209	8395940
CACNG2	calcium channel, voltage-dependent, gamma subunit 2	reserved	602911	
CACNG3	calcium channel, voltage-dependent, gamma subunit 3	reserved		
CACNG4	calcium channel, voltage-dependent, gamma subunit 4	17q24		10613843
CACNG5	calcium channel, voltage-dependent, gamma subunit 5	17q24		10613843

TABLE 1 (Continued)

Some ion channel genes of interest for EFS experiments				
Symbol	Full Name	Cytogenetic Location	MIM Number	PubMed ID
CACNG6	calcium channel, voltage-dependent, gamma subunit 6	19q13.4		11170751
CACNG7	calcium channel, voltage-dependent, gamma subunit 7	19q13.4		11170751
CACNG8	calcium channel, voltage-dependent, gamma subunit 8	19q13.4		11170751
KCNA1	potassium voltage-gated channel, shaker-related subfamily, member 1 (episodic ataxia with myokymia)	12p13	176260	1349297
KCNA1B	literature alias, see KCNAB1	-		
KCNA2	potassium voltage-gated channel, shaker-related subfamily, member 2	12	176262	
KCNA2B	literature alias, see KCNAB2	-		
KCNA3	potassium voltage-gated channel, shaker-related subfamily, member 3	1p13.3 or 13	176263	2251283
KCNA3B	literature alias, see KCNAB3	-		
KCNA4	potassium voltage-gated channel, shaker-related subfamily, member 4	11p14	176266	2263489
KCNA4L	potassium voltage-gated channel, shaker-related subfamily, member 4-like	11q14		8449523
KCNA5	potassium voltage-gated channel, shaker-related subfamily, member 5	12	176267	
KCNA6	potassium voltage-gated channel, shaker-related subfamily, member 6	reserved	176257	
KCNA7	potassium voltage-gated channel, shaker-related subfamily, member 7	19	176268	
KCNA8	literature alias, see KCNQ1	-		

TABLE 1 (Continued)

Some ion channel genes of interest for EFS experiments				
Symbol	Full Name	Cytogenetic Location	MIM Number	PubMed ID
KCNA9	symbol withdrawn, see KCNQ1	-		
KCNA10	potassium voltage-gated channel, shaker-related subfamily, member 10	reserved	602420	
KCNAB1	potassium voltage-gated channel, shaker-related subfamily, beta member 1	3q26.1	601141	8838324
KCNAB2	potassium voltage-gated channel, shaker-related subfamily, beta member 2	1p36.3	601142	8838324
KCNAB3	potassium voltage-gated channel, shaker-related subfamily, beta member 3	17p13.1	604111	9857044
KCNB1	potassium voltage-gated channel, Shab-related subfamily, member 1	20q13.2	600397	7774931
KCNB2	potassium voltage-gated channel, Shab-related subfamily, member 2	8		9612272
KCNC1	potassium voltage-gated channel, Shaw-related subfamily, member 1	11p15	176258	8449507
KCNC2	potassium voltage-gated channel, Shaw-related subfamily, member 2	12 and 19q13.4	176256	8111118
KCNC3	potassium voltage-gated channel, Shaw-related subfamily, member 3	19	176264	1740329
KCNC4	potassium voltage-gated channel, Shaw-related subfamily, member 4	1p21	176265	1920536
KCND1	potassium voltage-gated channel, Shal-related subfamily, member 1	Xp11.23-p11.3	300281	10729221
KCND2	potassium voltage-gated channel, Shal-related subfamily, member 2	7q31-32	605410	10551270
KCND3	potassium voltage-gated channel, Shal-related subfamily, member 3	1p13.2	605411	10942109
KCNE1	potassium voltage-gated channel, Isk-related family, member 1	21q22.1-q22.2	176261	8432548

TABLE 1 (Continued)

Some ion channel genes of interest for EFS experiments				
Symbol	Full Name	Cytogenetic Location	MIM Number	PubMed ID
KCNE1L	potassium voltage-gated channel, Isk-related family, member 1-like	Xq22.3	300328	10493825
KCNE2	potassium voltage-gated channel, Isk-related family, member 2	21q22.1	603796	10219239
KCNE3	potassium voltage-gated channel, Isk-related family, member 3	reserved	604433	10219239
KCNE4	potassium voltage-gated channel, Isk-related family, member 4	reserved		10219239
KCNF1	potassium voltage-gated channel, subfamily F, member 1	2p25	603787	9434767
KCNF2	literature alias, see KCNG2	-		
KCNF	symbol withdrawn, see KCNF1	-		
KCNG1	potassium voltage-gated channel, subfamily G, member 1	20q13	603788	9434767
KCNG2	potassium voltage-gated channel, subfamily G, member 2	18q22-18q23	605696	10551266
KCNG	symbol withdrawn, see KCNG1	-		
KCNH1	potassium voltage-gated channel, subfamily H (eag-related), member 1	1q32-41	603305	9738473
KCNH2	potassium voltage-gated channel, subfamily H (eag-related), member 2	7q35-q36	152427	7842012
KCNH3	potassium voltage-gated channel, subfamily H (eag-related), member 3	12q13	604527	10455180
KCNH4	potassium voltage-gated channel, subfamily H (eag-related), member 4	reserved	604528	10455180
KCNH5	potassium voltage-gated channel, subfamily H (eag-related), member 5	14	605716	9738473
KCNIP1	Kv channel interacting protein 1	reserved		10676964
KCNIP2	Kv channel-interacting protein 2	10		10676964

TABLE 1 (Continued)

Some ion channel genes of interest for EFS experiments				
Symbol	Full Name	Cytogenetic Location	MIM Number	PubMed ID
KCNIP3	literature alias, see CSEN	-		
KCNJ1	potassium inwardly-rectifying channel, subfamily J, member 1	11q24	600359	7680431
KCNJ2	potassium inwardly-rectifying channel, subfamily J, member 2	17q23.1-q24.2	600681	7696590
KCNJ3	potassium inwardly-rectifying channel, subfamily J, member 3	2q24.1	601534	8088798
KCNJ4	potassium inwardly-rectifying channel, subfamily J, member 4	22q13.1	600504	8016146
KCNJ5	potassium inwardly-rectifying channel, subfamily J, member 5	11q24	600734	
KCNJ6	potassium inwardly-rectifying channel, subfamily J, member 6	21q22.1	600877	7796919
KCNJ7	symbol withdrawn, see KCNJ6	-		
KCNJ8	potassium inwardly-rectifying channel, subfamily J, member 8	12p11.23	600935	8595887
KCNJ9	potassium inwardly-rectifying channel, subfamily J, member 9	1q21-1q23	600932	8575783
KCNJ10	potassium inwardly-rectifying channel, subfamily J, member 10	1q	602208	9367690
KCNJ11	potassium inwardly-rectifying channel, subfamily J, member 11	11p15.1	600937	7502040
KCNJ12	potassium inwardly-rectifying channel, subfamily J, member 12	17p11.1	602323	7859381
KCNJ13	potassium inwardly-rectifying channel, subfamily J, member 13	2q37	603208	9878260
KCNJ14	potassium inwardly-rectifying channel, subfamily J, member 14	19q13	603953	9592090

TABLE 1 (Continued)

Some ion channel genes of interest for EFS experiments				
Symbol	Full Name	Cytogenetic Location	MIM Number	PubMed ID
KCNJ15	potassium inwardly-rectifying channel, subfamily J, member 15	21q22.2	602106	9299242
KCNJ16	potassium inwardly-rectifying channel, subfamily J, member 16	17q23.1-q24.2	605722	11240146
KCNJN1	channel, subfamily J, inhibitor 1	17p11.2-p11.1	602604	8647284
KCNK1	potassium channel, subfamily K, member 1 (TWIK-1)	1q42-q43	601745	8661042
KCNK2	potassium channel, subfamily K, member 2 (TREK-1)	1q41	603219	9721223
KCNK3	potassium channel, subfamily K, member 3 (TASK-1)	2p23	603220	9312005
KCNK4	potassium inwardly-rectifying channel, subfamily K, member 4	11q13	605720	10767409
KCNK5	potassium channel, subfamily K, member 5 (TASK-2)	6p21	603493	9812978
KCNK6	potassium channel, subfamily K, member 6 (TWIK-2)	19q13.1	603939	10075682
KCNK7	potassium channel, subfamily K, member 7	11q13	603940	10206991
KCNK9	potassium channel, subfamily K, member 9 (TASK-3)	8	605874	10734076
KCNK10	potassium channel, subfamily K, member 10	reserved	605873	
KCNK12	potassium channel, subfamily K, member 12	2p22-2p21		
KCNK13	potassium channel, subfamily K, member 13	14q24.1-14q24.3		11060316

TABLE 1 (Continued)

Some ion channel genes of interest for EFS experiments				
Symbol	Full Name	Cytogenetic Location	MTM Number	PubMed ID
KCNK14	potassium channel, subfamily K, member 14	2p22-2p21		11060316
KCNK15	potassium channel, subfamily K, member 15	reserved		
KCNMA1	potassium large conductance calcium-activated channel, subfamily M, alpha member 1	10	600150	7987297
KCNMB1	potassium large conductance calcium-activated channel, subfamily M, beta member 1	5q34	603951	8799178
KCNMB2	symbol withdrawn, see KCNMB3	-		
KCNMB2	potassium large conductance calcium-activated channel, subfamily M, beta member 2	reserved	605214	10097176
KCNMB2L	symbol withdrawn, see KCNMB3L	-		
KCNMB3	potassium large conductance calcium-activated channel, subfamily M beta member 3	3q26.3-q27	605222	10585773
KCNMB3L	potassium large conductance calcium-activated channel, subfamily M, beta member 3-like	22q11		10585773
KCNMB4	potassium large conductance calcium-activated channel, subfamily M, beta member 4	reserved	605223	
KCNMBL	symbol withdrawn, see KCNMB3	-		
KCNMBLP	symbol withdrawn, see KCNMB3L	-		
KCNN1	potassium intermediate/small conductance calcium-activated channel, subfamily N, member 1	19p13.1	602982	8781233

TABLE 1 (Continued)

Some ion channel genes of interest for EFS experiments				
Symbol	Full Name	Cytogenetic Location	MIM Number	PubMed ID
KCNN2	potassium intermediate/small conductance calcium-activated channel, subfamily N, member 2	reserved	605879	
KCNN3	potassium intermediate/small conductance calcium-activated channel, subfamily N, member 3	22q11-q13.1	602983	9491810
KCNN4	potassium intermediate/small conductance calcium-activated channel, subfamily N, member 4	19q13.2	602754	9380751
KCNQ1	potassium voltage-gated channel, KQT-like subfamily, member 1	11p15.5	192500	8528244
KCNQ1OT1	KCNQ1 overlapping transcript 1	11p15.5	604115	10220444
KCNQ2	potassium voltage-gated channel, KQT-like subfamily, member 2	20q13.3-2 20q13.3	121200	9425895
KCNQ3	potassium voltage-gated channel, KQT-like subfamily, member 3	8q24	121201	9425900
KCNQ4	potassium voltage-gated channel, KQT-like subfamily, member 4	1p34	603537	10025409
KCNQ5	potassium voltage-gated channel, KQT-like subfamily, member 5	6q14		10787416
KCNS1	potassium voltage-gated channel, delayed-rectifier, subfamily S, member 1	reserved	602905	9305895
KCNS2	potassium voltage-gated channel, delayed-rectifier, subfamily S, member 2	8q22	602906	9305895
KCNS3	potassium voltage-gated channel, delayed-rectifier, subfamily S, member 3	reserved	603888	10484328

PCR reactions can be carried out with a variety of thermostable enzymes including but not limited to AmpliTaq, AmpliTaq Gold, or Vent polymerase. For AmpliTaq, reactions can be carried out in 10 mM Tris-Cl, pH 8.3, 2.0 mM MgCl₂, 200 μ M of each dNTP, 50 mM KCl, 0.2 μ M of each primer, 10 ng of DNA
5 template, 0.05 units/ μ l of AmpliTaq. The reactions are heated at 95°C for 3 minutes and then cycled 35 times using suitable cycling parameters, including, but not limited to, 95°C, 20 seconds, 62°C, 20 seconds, 72°C, 3 minutes. In addition to these conditions, a variety of suitable PCR protocols can be found in PCR Primer, A Laboratory Manual, edited by C.W. Dieffenbach and G.S. Dveksler, 1995, Cold
10 Spring Harbor Laboratory Press; or PCR Protocols: A Guide to Methods and Applications, Michael et al., eds., 1990, Academic Press.

It is desirable to sequence the DNA encoding voltage-gated ion channels obtained by the herein-described methods, in order to verify that the desired voltage-gated ion channel has in fact been obtained and that no unexpected changes
15 have been introduced into its sequence by the PCR reactions. The DNA can be cloned into suitable cloning vectors or expression vectors, *e.g.*, the mammalian expression vector pcDNA3.1 (Invitrogen, San Diego, CA) or other expression vectors known in the art or described herein.

A variety of expression vectors can be used to recombinantly express
20 DNA encoding voltage-gated ion channels for use in the present invention. Commercially available expression vectors which are suitable include, but are not limited to, pMC1neo (Stratagene), pSG5 (Stratagene), pcDNAI and pcDNAIamp, pcDNA3, pcDNA3.1, pCR3.1 (Invitrogen, San Diego, CA), EBO-pSV2-neo (ATCC 37593), pBPV-1(8-2) (ATCC 37110), pdBPV-MMTneo(342-12) (ATCC 37224),
25 pRSVgpt (ATCC 37199), pRSVneo (ATCC 37198), pCI.neo (Promega), pTRE (Clontech, Palo Alto, CA), pV1Jneo, pIRESneo (Clontech, Palo Alto, CA), pCEP4 (Invitrogen, San Diego, CA), pSC11, and pSV2-dhfr (ATCC 37146). The choice of vector will depend upon cell type in which it is desired to express the voltage-gated ion channels, as well as on the level of expression desired, and the like.

30 The expression vectors can be used to transiently express or stably express the voltage-gated ion channels. The transient expression or stable expression of transfected DNA is well known in the art. See, *e.g.*, Ausubel et al., 1995, "Introduction of DNA into mammalian cells," in Current Protocols in Molecular Biology, sections 9.5.1-9.5.6 (John Wiley & Sons, Inc.).

As an alternative to the above-described PCR methods, cDNA clones encoding ion channels can be isolated from cDNA libraries using as a probe oligonucleotides specific for the desired voltage-gated ion channels and methods well known in the art for screening cDNA libraries with oligonucleotide probes. Such methods are described in, *e.g.*, Sambrook *et al.*, 1989, *Molecular Cloning: A Laboratory Manual*; Cold Spring Harbor Laboratory, Cold Spring Harbor, New York; Glover, D.M. (ed.), 1985, *DNA Cloning: A Practical Approach*, MRL Press, Ltd., Oxford, U.K., Vol. I, II. Oligonucleotides that are specific for particular voltage-gated ion channels and that can be used to screen cDNA libraries can be readily designed based upon the known DNA sequences of the voltage-gated ion channels and can be synthesized by methods well-known in the art.

The present invention also provides apparatuses for use with the methods disclosed herein. For example, the present invention provides a multiwell tissue culture plate where a plurality of the wells of the plate contain a pair of electrodes disposed such that when a preselected voltage is applied across the electrodes the transmembrane potential of cells within the wells is altered.

In certain embodiments, the multiwell tissue culture plate contains one of the pair of electrodes on the bottom of the wells and the other of the pair of electrodes on the side of the wells. This embodiment is depicted in Figure 2B.

In other embodiments, the multiwell tissue culture plate contains both of the pair of electrodes on the bottom of the wells. This embodiment is depicted in Figure 2C.

In other embodiments of the multiwell tissue culture plate, one of the pair of electrodes is a layer of conductive material that forms the bottom of the wells and the other of the pair of electrodes enters the wells from above. This embodiment is depicted in Figures 7, 12, and 16.

In other embodiments of the multiwell tissue culture plate, both of the pair of electrodes are embedded in an insulator and enter the wells from above. This embodiment is depicted in Figures 9 and 10.

In other embodiments of the multiwell tissue culture plate, the electrode that enters the wells from above has a central conductive material portion that is surrounded by an insulator. This embodiment is depicted in Figure 8.

In other embodiments of the multiwell tissue culture plate, one of the pair of electrodes forms the bottom of the wells and the other of the pair of electrodes enters the wells from above. This embodiment is depicted in Figures 7 and 10.

5 In other embodiments of the multiwell tissue culture plate, the pairs of electrodes form an alternating pattern of positive and negative electrodes in the wells. This embodiment is depicted in Figure 16.

10 In other embodiments of the multiwell tissue culture plate, the layer of conductive material that forms the bottom of the wells is a layer of indium tin oxide that overlays a glass substrate. Preferably, the layer of conductive material and the glass substrate are transparent.

In other embodiments of the multiwell tissue culture plate, a plurality of the wells of the plate contain interdigitating electrodes. This embodiment is depicted in Figures 3 and 5.

15 The present invention provides a multiwell tissue culture plate where: the bottom of the wells is a filter membrane upon which cells can be grown;

the wells are located in a trough that can contain fluid;

the trough contains a first electrode;

a second electrode enters the wells from above;

20 where the first and second electrodes are so disposed that when a preselected voltage is applied across the electrodes the transmembrane potential of cells within the wells is altered. This embodiment is depicted in Figure 8.

25 The present invention also provides a combination of the multiwell tissue culture plates disclosed herein and a fluorescent imager where the multiwell tissue culture plate and the fluorescent imager are positioned relative to one another such that the fluorescent imager can obtain fluorescent readings from the wells of the multiwell tissue culture plate.

The present invention also provides a combination of a top substrate and a bottom substrate where the top and bottom substrates each contain:

30 a plurality of virtual wells; and

a layer of conductive material that forms the bottoms of the virtual wells; where the layers of conductive material in the top and bottom substrates are connected to a pulse generator such that the layers of conductive material function as electrodes such that when a preselected voltage is applied across the electrodes the

transmembrane potential of cells within the virtual wells is altered. Such a combination is depicted in Figures 6 and 13.

The present invention also provides a substrate having square or rectangular wells formed by a plurality of generally parallel positive and negative electrodes and a plurality of spacers arranged generally at right angles to the electrodes, where:
one wall of the wells is formed by a positive electrode and the opposite wall of the well is formed by a negative electrode;
the spacers form the walls of the wells that are at right angles to the walls formed by the electrodes;
where the electrodes are so disposed that when a preselected voltage is applied across the electrodes the transmembrane potential of cells within the wells is altered. Such a substrate is depicted in Figure 1.

An example of another embodiment of the present invention comprises:
a substrate having an upper surface upon which are present at least 10^3 living eukaryotic cells which have a voltage-gated ion channel of interest in their plasma membranes;
a plurality of positive electrodes and a plurality of negative electrodes positioned either on or near the substrate such that when a voltage is applied through the positive and negative electrodes the transmembrane potential of the cells is altered;
at least one substance that is suspected of being an activator or an inhibitor of the voltage-gated ion channel;
where the cells contain a fluorescent indicator compound.

An example of another embodiment of the present invention comprises:
a multiwell tissue culture plate having a plurality of wells in which are present at least 10^3 living eukaryotic cells per well of the plurality which cells have a voltage-gated ion channel of interest in their plasma membranes;
a plurality of positive electrodes and a plurality of negative electrodes positioned such that when a preselected voltage is applied through the positive and negative electrodes, the transmembrane potential of the cells is altered;

at least one substance that is suspected of being an activator or an inhibitor of the voltage-gated ion channel in at least one of the plurality of the wells; where the cells contain a fluorescent indicator compound or a voltage sensitive membrane dye.

5

The following non-limiting examples are presented to better illustrate the invention.

Example 1

10

In Figure 24, a preferred system for conducting high throughput screening using EFS stimulation is shown. The system consist of a computer 2402 that comprises an arbitrary waveform generator card 2404 electronically associated with the computer 2402. Custom software was written on the computer 2402 which causes the arbitrary generator card 2404 to generate a pulse voltage waveform (2406) of the appropriate electrical stimulus. The voltage waveform (2406) is applied to the input of eight constant current amplifiers 2408. Each constant current amplifier 2408 services a row on the 96-well sample filter plate 2410. The outputs from the amplifiers 2412 pass through the contacts of electrical relays 2414 allowing the current pulse to be applied to the electrodes 2416.

20

The waveform generator card 2404 also generates a 7-bit binary transistor-transistor logic TTL value (2418) that represents the address of the well to be excited by the stimulus. In addition, a trigger pulse 2420 is generated. Microprocessor controller 2422, waits for the trigger pulse 2420, interprets the binary value (2418) and then switches on the appropriate relay 2414 which then directs the constant current pulse (2424) to the particular electrode 2416 or electrodes, via electrode connecting wire(s) 2417 in the sample well 2426. Current flows from the amplifier's output (2424), through the relay contact 2414 through the electrode 2416 the liquid in the well 2428, through the well's membrane 2430 and returns via fluid 2432 beneath the membrane 2430 and a return wire 2434. One large common current return trough 2436 services

30

all 96-electrodes. Other arrangements are possible where each sample well has its own isolated current return trough and return wire. See Example 2 below.

The current return trough 2436 beneath the membranes 2430 has a clear glass bottom 2438 that permits excitation light (2440) from a light source 2442 to pass through the glass bottom 2438, through the transparent membrane 2430 and illuminate cells 2444 adhered to the membrane 2430. Fluorescent light (2446) from the cells 2444 returns back through the membrane 2430 and the glass bottom 2438 entering into the detector 2448. Suitable detectors include those described *supra*. The preferred detector is the FLIPR (Molecular Devices) fluorescence imager ^{on the VIPR (Auto ra Biosciences)}

When the pulse sequence is completed, the microprocessor controller 2422 switches off the relays 2414 isolating the constant current amplifiers' pulses (2424) from the electrodes 2416.

Turning to Figures 25 and 26, Figure 25 represents a photograph of an electrode head 2500 embodiment comprising top electrodes 2516 and first electrode connecting wires 2517. The electrode head comprises a ground contact rod 2510. Figure 26 represents a photograph of a trough embodiment 2600 for use in conjunction with the electrode head 2500 embodiment shown in Figure 25. The trough 2600 comprises bracing posts 2610 to assist in aligning and attachment of the electrode head through apertures 2520 in the electrode head 2500 (see Figure 28). A bottom electrode wire (hidden) is positioned in the trough which when submerged in the salt/buffer solution, upon assembly of the EFS system (see Figure 28) acts as bottom electrode for each of the wells. The bottom electrode wire is in electrical communication with a return connection wire 2620 at position 2630. The return connection wire is secured to the ground contact rod 2510 upon assembly of the EFS system. The trough 2600 also comprises a transparent bottom portion 2640 preferably made of glass.

Figure 27 represents a photograph of the trough embodiment 2600 wherein a MultiscreenTM-Black CM 96 wellplate 2700, with 96 wells 2710, is positioned in the

trough 2600. Information concerning Millipore's multiscreen plates and biopore membranes is found, e.g., at <http://www.millipore.com/catalogue.nsf/docs/C7781> and <http://www.millipore.com/publications.nsf/docs/tn062>.

5 Figure 28 is a photograph of the assembled EFS system 2800 comprising the trough 2600 with well plate 2700 in place. The electrode head 2500 is secured to the top of the trough 2600 such that the electrodes 2416 are inserted into the wells 2710, one electrode per well. The electrode head 2500 is secured down onto bracing posts 2610 (hidden) by fasteners 2810. The fasteners are preferable threaded nuts.

10 Preferably, prior to assembly, each well 2710 (hidden) has been loaded with cells which have been cultured to canvas the bottom of the wells 2710 (hidden). After cells have been cultured under standard and known conditions, and before assembly of the EFS unit 2800, each well is preferably washed to remove cell media and then loaded with the predetermined buffer solution as discussed above.

15 Figure 29 shows a graphical representation of data obtained from an embodiment of the invention similar to that depicted in Figure 28. The data represent a membrane potential change in HEK293 cells that have been transfected to express human PN1 voltage-gated sodium channel. Each plot represents a row (12wells) A-H

20 of a 96-well plate. Each column of the 96-well plate data was acquired for 15 seconds on a VIPR™. Stimulation pulse protocol was applied during the data acquisition as follows; 2s baseline was followed with a 2ms square pulse, Amplitude = 20mA, Frequency = 10 Hz, Duration = 5s. Those skilled in the art will readily appreciate, in view of the teachings herein, that the subject system may generate a pulse between

25 1 μ s to 1s. Preferably, the pulse generated is between about 0.1ms and about 100ms.

 Figure 30 is a bar graph representation of the peak ratio change of data depicted in Figure 29. 1 μ M TTX a specific and potent blocker of tetrodotoxin (TTX)-

30 sensitive voltage-gated sodium channels is present in wells E1, F1, G1, H1, A12, B12,

C12 and D12. In addition well A11 contains an internal standard for blocking TTX-sensitive voltage-gated sodium channels. Z-score is a measure of the difference in the ^{of the uninhibited and inhibited signals} uninhibited and inhibited signal divided by the sum of the standard deviations.

5 Figure 31 shows the effects of increasing concentrations of TTX (upper panel) and of Compound A (lower panel) on the EFS-stimulated depolarization signal in HEK293/PN1 cells. The IC₅₀s obtained in these experiments are comparable to those obtained through other techniques. The high Hill coefficients, nH, result from the threshold nature of the stimulation protocol ^{and the nonlinear relation between} _{ion channel activity and} _{membrane potential.}

10 Example 2

 Figure 32 represents a photograph of an EFS embodiment 3200 pertaining to an alternative EFS system configuration. The electrode head 2500 is similar to that described above in Figure 25. However, the configurations of the electrodes, wells and trough are configured differently to further isolate the electrical fields. This reduces cross-talk and interference between wells. For this embodiment, the inventors have adapted Millipore's MultiscreenTM Caco-2 Assay System for use as a EFS system. Information concerning the MultiscreenTM Caco-2 Assay System can be found at <http://www.millipore.com/publications.nsf/docs/PF1780EN00>. The standard commercially available Caco-2 plate system comprises two plates: a membrane-bottom cell growth plate and a 96-well receiver tray. One of the unique characteristics of the Caco-2 system is that it each well has an individual corresponding trough that is accessed basolaterally to each well. Therefore, it supplants the need for a common trough into which all of the wells sit. According to this embodiment, the top electrodes 2516 are disposed into each of the wells in the membrane-bottom cell growth plate (hidden). To establish the bottom electrode for each well, a conductive electrode plate 3220 is provided. The conductive electrode plate 3220 comprise a series of well apertures 3230, providing access of the top electrodes 2410 into the individual wells during assembly. The conductive electrode

15

20

25

30

plate 3220 also comprises a series of conductive pins (hidden) secured thereto and extending downward at positions 3240. These conductive pins are inserted through the basolateral access port of the membrane-bottom cell growth plate (not shown).

5 Figure 33 is a depiction of the bottom of the conductive electrode plate 3220 and shows the conductive pins 3310, which are extending out of the page toward the reader. Figure 34 shows a side-view of the conductive electrode plate 3220 properly positioned atop of the membrane-bottom cell growth plate 3410 and 96 well receiver tray 3420. When the electrode conductive plate 3220 is properly positioned on top of
10 the membrane-bottom cell growth plate 3410, the conductive pins 3310 are inserted through the basolateral access port (not shown) into the individual trough area (not shown) of the 96 well receiver tray. When the individual trough area is filled with the appropriate solution it contacts the bottom of each well and individual pin. Therefore, when the well and trough area are filled with solution, current may flow from the top
15 electrode to the bottom electrode during operation. Figure 35 is a side-view of the assembled EFS system. The assembled system comprises the membrane-bottom cell growth plate 3410 positioned in the 96 well tray 3420. The electrode plate 3220 is mounted on top of the membrane bottom well plate 3410. The electrode head 2500 is shown mounted on top of the electrode plate 3220.

20

One clear advantage to the EFS systems described in Examples 1 and 2 above, and elsewhere in the present application, is the ability to generate a uniform field across the cells, as opposed to tangential to the cells. Generating an electrical field across the cells is made possible by the novel "top to bottom" placement of the
25 electrodes in a multiwell format.

Example 3

Figure 36 shows a novel electrode embodiment 3600. Figure 36A depicts an
30 expanded view of the electrode 3600. The electrode 3600 comprises two parallel

plates 3610 and 3630 with a low dielectric plate or disc 3620 between them.

Optionally, the electrode may be coated with an insulating material. Potential advantages of this design are that special multiwell plates are not required, i.e., any plate that the cells will stick to and that the stimulation and emission light will pass through may be used. There is no filter in the well that may absorb compound or pass compound during long incubations. In the case of the coated electrode, very little current is used and ohmic heating is diminished, even for dc current and even for extended periods of stimulation. The capacitance current is low enough that this advantage applies to ac current as well. The sealed electrode permits placement very close to the cell layer for more uniform stimulation.

Not to be bound by any theory, it is believed that the more uniform the electrical field presented to the cells is, a more accurate indication of potential modulation to the cells will be achieved. In other words, the more uniform the electrical field is, the potential modulation as observed by any of the methods presented herein, e.g., fluorescence, will more directly correlate to actual modulation of ion channels in the cell membrane, and less correlate with background noise in the system caused by cross-interference, cross-illumination, dye effects, dye leaching or any other interference in the system. One way to increase the uniformity of the electrical field applied to the cells is to present one or more of the electrodes in close proximity to, or in contact with, the cells. However, this can affect the cells in deleterious ways leading to failure in the system. Some of the problems associated with close proximity or contact of the electrode(s) to the cells are caused by, for example, ohmic heating, oxidation and formation of bubbles on the electrode. The embodiments of the present invention as taught in Figures 8, 11, 24-28 and 32-35 are particularly preferred because they achieve a uniform electrical field across the cells without putting the electrodes in contact with or close proximity to the cells. Furthermore, the novel electrode design shown in Figure 36 achieves a uniform electrical field, by allowing close proximity of the electrode to the cells, without creating the problems of ohmic heating, oxidation, or bubbling of the cells.

It is believed that the subject EFS system embodiments produce substantially uniform fields, where the one or more electrical fields vary over an area of observation by no more than about 30% from the mean electrical field at any one
5 time. Percentages are determined by measurements in two dimensions; or preferably, variation is calculated in three dimensions. In a more preferred embodiment, the one or more electrical fields vary over an area of observation by no more than about 15 % from the mean electrical field at any one time. In an even more preferred embodiment, the one or more electrical fields vary over an area of observation by no
10 more than 10 % from the mean electrical field at any one time. In an optimal embodiment, the variation is no more than 5 % from the mean.

The similarity to a capacitor is obvious, but the low dielectric 3620 between the plates 3610 and 3630 reduces the amount of current required to initially charge the
15 plates with only a miniscule current required to maintain the charge between the plates. An external electric field is generated that can be used to depolarize the cells. The external electric field density is reduced by a high dielectric between the plates as is used with an authentic capacitor and is maximal with a low dielectric such as teflon or mylar or no dielectric. The external field density is further enhanced by placing the
20 plates very close together, but the optimal separation may be determined empirically.

Figure 36B shows an embodiment comprising a concurrent lead design. The concurrent lead comprises an internal wire 3655 and an external wire 3650. The internal wire passes through the top plate 3610 and dielectric plate 3620 and is
25 attached or integral to the bottom plate 3630. The external wire is attached or integral to the top plate 3610. Those skilled in the art will recognize that the foregoing arrangement of the leads may be reversed. Figure 36C shows an embodiment comprising edge leads 3660 and 3665. Edge lead 3660 is attached or integral to top plate 3610 and edge lead 3665 is attached or integral to bottom plate 3630.

Some of the embodiments of the subject invention include the following:

5

A method of characterizing the biological activity of a candidate compound comprising.

exposing one or more cells to said compound; repetitively exposing said one or more cells to one or more electric fields so as to effect a controlled change in
10 transmembrane potential of said one or more cells; and monitoring, without using a patch clamp, changes in the transmembrane potential of said one or more cells.

The above method, where the monitoring comprises detecting fluorescence emission from an area of observation containing said one or more cells.

The above method, where the electric fields are biphasic.

15 The above method, additionally comprising limiting spatial variation in electric field intensity so as to minimize irreversible cell electroporation.

The above method, where one or more electrical fields may cause an ion channel of interest to cycle between different voltage dependent states.

20 The above method, where the one or more electrical fields cause an ion channel of interest to open.

The above method, where the one or more electrical fields cause an ion channel of interest to be released from inactivation.

25 The above method, where the one or more cells comprise a voltage sensor selected from the group consisting of a FRET based voltage sensor, an electrochromic transmembrane potential dye, a transmembrane potential redistribution dye, an ion sensitive fluorescent or luminescent molecule and a radioactive ion.

The above method, where the one or more cells comprise a voltage regulated ion channel.

The above method, where the voltage regulated ion channel is selected from the group consisting of a potassium channel, a calcium channel, a chloride channel and a sodium channel.

5 The above method, where the electric field exhibits limited spatial variation in intensity in the area of observation of less than about 25% from a mean intensity in that area.

The above method, where the one or more electrical fields varies over an area of observation by no more than about 15 % from the mean electrical field at any one time.

10 The above method, where the one or more electrical fields varies over an area of observation by no more than about 5 % from the mean electrical field at any one time.

The above method, where the one or more electrical fields comprises stimulation with either a square wave-form, a sinusoidal wave-form or a saw tooth wave-form.

15 The above method, where the one or more electrical fields have an amplitude within the range of about 10 V/cm to about 100 V/cm.

The above method, where the one or more electrical fields have an amplitude within the range of about 20 V/cm to about 80 V/cm.

20 The above method, where the one or more electrical fields are repeated at a frequency of stimulation that is greater than or equal to the reciprocal of the transmembrane time constant of said one or more cells.

The above method, where the one or more electrical fields are repeated at a frequency of stimulation within the range of zero to 1 kHz.

The above method, where the one or more electrical fields have a pulse duration within the range of about 100 microseconds to about 20 milliseconds.

25 The above method, where the transmembrane potential is developed across the plasma membrane of said one or more cells.

A method of assaying the biochemical activity of a compound against a target ion channel comprising.

30 selecting a cell line having a normal resting transmembrane potential corresponding to a selected voltage dependent state of said target ion channel; expressing said target

ion channel in a population of cells of said selected cell line; exposing said population of cells to said compound; repetitively exposing said population of cells to one or more electric fields so as to effect a controlled change in transmembrane potential of said one or more cells; and monitoring changes in the transmembrane potential of said
5 one or more cells.

The above method, where the target ion channel is exogenously expressed in said cell line.

The above method, where the cell line is transfected with nucleic acid encoding said target ion channel.

10 The above method, where the cell line expresses insignificant levels of other ion channels.

The above method, where the cell line is selected from the group consisting of CUL, LTK(-), and CHO-M.

15 The above method, where the target ion channel is a sodium channel, and wherein said population of cells is selected from the group consisting of CHL cells, LTK(-) cells, and CHO-K1 cells.

The above method, where the target ion channel is a sodium channel, and wherein said population of cells is selected from the group consisting of HEK-293 cells, RBL cells, F11 cells, and HL5 cells.

20 The above method, where the target ion channel is a potassium channel, and wherein said population of cells is selected from the group consisting of CHL cells, LTK(-) cells, and CHO-K1 cells.

The above method, where the target ion channel is a calcium channel, and wherein said population of cells is selected from the group consisting of CHL cells,
25 LTK(-) cells, and CHO-K1 cells.

A method of assaying ion channel activity comprising.

exposing at least one cell to a plurality of electric field pulses so as to create a controlled change in transmembrane potential and so as to activate an ion channel of interest; and detecting ion channel activity by detecting one or more changes in
30 transmembrane potential without using a patch clamp.

The above method, where the at least one cell comprises a voltage sensor selected from the group consisting of a FRET based voltage sensor, an electrochromic transmembrane potential dye, a transmembrane potential redistribution dye, an ion sensitive fluorescent or luminescent molecule and a radioactive ion.

- 5 The above method, where the voltage sensor comprises a FRET based voltage sensor.

The above method, where the ion channel of interest is a voltage regulated ion channel.

- The above method, where the plurality of electric field pulses cause said ion
10 channel of interest to cycle between different voltage dependent states.

The above method, where the at least one cell is an eukaryotic cell.

The above method, where the at least one cell is a non-excitable cell.

The above method, where the at least one cell is a prokaryotic cell.

The above method, where the at least one cell is a tissue culture cell.

- 15 The above method, where the at least one cell is a primary cell line.

The above method, where the at least one cell is part of an intact living organism.

A method of assaying ion channel activity comprising.

- expressing a selected target ion channel in at least one cell; expressing a selected
counter ion channel in said at least one cell; exposing said at least one cell to a
20 plurality of electric field pulses so as to create a controlled change in transmembrane
potential and so as to activate said counter ion channel; and monitoring the
transmembrane potential of said at least one cell.

The above method, where a transmembrane potential change is detected when said
ion channel of interest is blocked.

- 25 The above method, where the ion channel of interest comprises a ligand gated ion
channel.

The above method, where the counter channel comprises a sodium channel.

- A method of modifying the transmembrane potential of a cell comprising
repetitively applying biphasic electric field pulses to said cell, wherein said pulses
30 have a maximum amplitude of less than approximately 90 V/cm, wherein said pulses

are applied at a rate of at least about 1 per second, and wherein the total duration of each pulse is at least about 1 millisecond.

The above method, where the maximum amplitude is approximately 20 to 40 V/cm.

The above method, where the pulse duration is approximately 2 to 10 milliseconds
5 per phase.

The above method, where the pulses are applied at a rate of approximately 20 to 100 pulses per second.

A method of characterizing the biological activity of a candidate compound comprising.

10 placing one or more cells into an area of observation in a sample well; exposing said one or more cells to said compound; repetitively exposing said one or more cells to a series of biphasic electric fields at a rate of approximately 20 to 100 pulses per second, wherein said electric fields exhibit limited spatial variation in intensity in the area of observation of less than about 25% from a mean intensity in that area, and
15 wherein said electric fields produce a controlled change in transmembrane potential of said one or more cells; and monitoring changes in the transmembrane potential of said one or more cells by detecting fluorescence emission of a FRET based voltage sensor from, an area of observation containing said one or more cells.

The above method, where the one or more electrical fields cause an ion channel of
20 interest to open.

The above method, where the one or more electrical fields cause an ion channel of interest to be released from inactivation.

The above method, where the one or more cells comprise a voltage regulated ion channel.

25 The above method, where the voltage regulated ion channel is selected from the group consisting of a potassium channel, a calcium channel, a chloride channel and a sodium channel.

The above method, where the one or more electrical fields likely vary over an area of observation by no more than about 15 % from the mean electrical field at any one
30 time.

The above method, where the one or more electrical fields varies over an area of observation by no more than about 5 % from the mean electrical field at any one time.

The above method, where the one or more electrical fields are selected from a square wave-form, a sinusoidal wave-form or a saw tooth wave-form.

5 A high throughput screening system comprising.

 a plurality of wells having a high transmittance portion through which cells present in said wells are optically observable in an area of observation; two electrodes in each of said plurality of wells; an optical detector configured to detect light emanating from said wells through said high transmittance portion; a power supply connected to
10 said electrodes; wherein said power supply and said electrodes are configured to apply a series of electric fields to cells within said area of observation, said electric fields having a spatial variation of less than about 25% of a mean field intensity within said area of observation, said electric fields being effective to controllably alter the transmembrane potential of a portion of said cells; a data processing unit configured
15 to interpret said light emanating from said wells, through said high transmittance portion as ion channel activity resulting from said transmembrane potential alterations.

 The above high throughput screening system, where the plurality of wells are located in a multiwell plate.

20 The above high throughput screening system, where the high transmittance portion is made from a material selected from the group consisting of glass, quartz, cycloolefin, Aclar, polypropylene, polyethylene and polystyrene.

 The above high throughput screening system, where the high transmittance portion exhibits less fluorescence when excited with UV light in the range of 250 nm to 400
25 nm than polystyrene.

 The above high throughput screening system, where the electrodes are located in a well of said plurality of wells.

 The above high throughput screening system, where the electrodes are located in a bottom layer of said plurality of wells.

The above high throughput screening system, where the multiwell plate comprises up to 96 wells.

The above high throughput screening system, where the multiwell plate comprises greater than 96 wells.

- 5 The above high throughput screening system, where the multiwell plate comprises greater than 384 wells.

The above high throughput screening system, where the electrodes are made of a material selected from the group consisting of gold, platinum, palladium, chromium, molybdenum, iridium, tungsten, tantalum and titanium.

- 10 The above high throughput screening system, where the multiwell plate comprises optically opaque materials or pigments to reduce the transmission of light.

The above high throughput screening system, where the electrodes are separated by a gap within the range of about 1 to 4 mm.

- 15 The above high throughput screening system, where the electrodes are separated by a gap within the range of about 0.1 to 1 mm.

1.0 The above high throughput screening system, where the electrodes are separated by a gap within the range of about 0.01 to 0.1 mm.

- 20 The above high throughput screening system, where the electrodes are charged to create an electrical field intensity of between 5 to 100 V/cm across said gap, and wherein the total charge transferred across the surface area of the electrically conductive material, in fluidic connection with the interior of the well is less than or equal to 100 μ C/mm².

- 25 The above high throughput screening system, where the plurality of wells further comprise an insulator orientated and configured so as to create an area of observation within said well in which the electrical field intensity varies by no more than 10 % from the mean electrical field intensity when said at least two strips of electrically conductive material are charged to create an electrical field intensity of between 5 to 100 V/cm across said gap, and wherein the total charge transferred across the surface area of the electrically conductive material, in fluidic connection with the interior of
30 the well is less than or equal to 100 μ C/mm².

The above high throughput screening system, where the plurality of wells further comprise at least two satellite electrical conductors.

A high throughput screening system comprising.

sample wells; liquid handling stations for adding reagents and/or cells to said
5 sample wells; and means for controlling the transmembrane potential of cells in said sample wells so as to selectively cause ion channel activity.

means for optically monitoring changes in said transmembrane potential.

The above high throughput screening system, where the means comprises electrodes configured to create an electric field having a spatial variation of less than
10 about 25% of a mean field intensity within an area of observation.

The above high throughput screening system, where the means for controlling the transmembrane potential comprise an electrode array assembly.

The above high throughput screening system, where the electrode assembly array comprises 8 electrode assemblies.

15 The above high throughput screening system, where the electrode assembly array comprises 96 electrode assemblies.

The above -high throughput screening system, where the electrode assembly array comprises greater than 96 electrode assemblies.

The above high throughput screening system, where the system further comprises
20 means for retractably moving said electrode assembly into and out of the wells of a multiwell plate.

The above high throughput screening system, where the means for controlling the transmembrane potential comprises electrical conductors with two substantially parallel planar surfaces.

25 The above high throughput screening system, where the electrical conductors are separated by a gap within the range of 1 to 4 mm.

The above high throughput screening system, where the electrical conductors are separated5 by a gap within the range of 0. 1 to 1 mm.

The above high throughput: screening system, where the electrical conductors
30 further comprise a first insulator.

The above high throughput screening system, where the first insulator comprises two planar surfaces orientated perpendicular to said substantially parallel planar surfaces of said electrical conductors and substantially parallel with respect to each other.

5 The above high throughput: screening system, where the electrical conductors further comprise a second insulator attached to said at least two electrical conductors, wherein said second insulator is interposed in said gap between said at least two electrical conductors to define the depth of said aqueous solution between said at least two electrical conductors.

10 The above high throughput: screening system, where the first insulator is composed of allow fluorescence material, wherein. said low fluorescence material exhibits less fluorescence when excited with UV light in the range 250 nm to 400 nm than polystyrene of comparable size.

The above high throughput screening system, where the second insulator is
15 composed of a low fluorescence material, wherein said low fluorescence material exhibits less fluorescence when excited with UV light in the range 250 nm to 400 nm than polystyrene of comparable size.

The above high throughput screening system, where the first insulator comprises an insulator selected from the group consisting of plastic, glass and ceramic.

20 The above high throughput screening system, where the plastic is selected from the group consisting of nylon, polystyrene, Teflon (tetrafluoroethylene), polypropylene, polyethylene, poly-vinyl chloride, and cycloolefin.

The above high throughput screening system, where the electrical conductors comprise a conductor selected from the group consisting of gold, platinum, titanium,
25 tungsten, molybdenum, iridium, vanadium, Nb, Ta, stainless steel and graphite.

The above high throughput screening system, where the electrical conductors comprise a surface treatment to reduce electrolysis.

The above high throughput screening system, where the surface treatment to reduce electrolysis comprises platinum black, gold black, iridium/iridium oxide,
30 titanium/titanium nitride or polypyrrole films.

The above high throughput screening system, where the electrical field intensity varies by no more than 10 % from the mean electrical field intensity when said at least two electrical conductors are charged to create an electrical field intensity of between 5 to 100 V/cm across said gap, wherein the total charge transferred across the surface area of the electrical conductors in contact with said aqueous solution is less than or
5 equal to 1 00 $\mu\text{C}/\text{mm}^2$.

The above high throughput screening system, where the electrical field intensity varies by no more than 5% from the mean electrical field intensity when said at least two electrical conductors are charged to create an electrical field intensity of between
10 5 to 100 V/cm across said gap, wherein the total charge transferred across the surface area of the electrical conductors in contact with said aqueous solution is less than or equal to 100 $\mu\text{C}/\text{mm}^2$.

A method of screening a plurality of drug candidate compounds against a target ion channel comprising.
15 expressing said target ion channel in a population of host cells; placing a plurality of said host cells into each of a plurality of sample wells; adding a candidate drug compound to at least: one of said plurality of sample wells; and modulating the transmembrane potential of host cells in said plurality of sample wells with a repetitive application of electric fields so as to set said transmembrane potential to a
20 level corresponding to a pre-selected voltage dependent state of said target ion channel.

The above method, additionally comprising selecting a host: cell line having a normal resting transmembrane potential corresponding to a second pre-selected voltage dependent state of said target ion channel.

25 The above method, where the electric fields are biphasic.

The above method, where electric fields cause an ion channel of interest to cycle between different voltage dependent states.

The above method, where the electric fields cause an ion channel of interest to open.

The above method, where the electric fields cause an ion channel of interest to be released from inactivation.

The above method, where the one or more cells comprise a voltage sensor selected from the group consisting of a FRET based voltage sensor, an electrochromic
5 transmembrane potential dye, a transmembrane potential redistribution dye, an ion sensitive fluorescent or luminescent molecule and a radioactive ion.

The above method, where the target ion channel is selected from the group consisting of a potassium channel, a calcium channel, a chloride channel and a sodium channel.

10 The above method, where the one or more electrical fields comprises stimulation with either a square wave-form, a sinusoidal wave-form or a saw tooth wave-form.

The above method, where the one or more electrical fields have an amplitude within the range of about 10 V/cm to about 100 V/cm.

The above method, where the one or more electrical fields have an amplitude
15 within the range of about 20 V/cm to, about 80 V/cm.

An assay plate and electrode assembly comprising at least one sample well having electrodes placed therein, wherein said electrodes are positioned with respect to the bottom surface of the well to provide an electric field adjacent to said bottom surface that varies by less than about 10% from a mean field intensity over at least about 20%
20 of the surface area of said bottom surface.

The above assembly, where the electrodes comprise plate electrodes extending down into said well such that bottom ends of said electrodes are adjacent to but not in contact with said bottom surface.

The above assembly, comprising two electrodes per sample well. The above
25 assembly, comprising more than two electrodes per sample well.

The above assembly, where the electrodes are plated onto said bottom surface of said well. The above assembly, where the bottom surface comprises a high optical transmittance portion.

The above assembly, where the high transmittance portion is made from a material selected from the group consisting of glass, quartz, cycloolefin, Aclar, polypropylene, polyethylene and polystyrene.

5 The above assembly, where the high transmittance portion exhibits less fluorescence when excited with UV light in the range of 250 nm to 400 nm than polystyrene.

The above assembly, where the electrodes are located in a wall of said plurality of wells.

The above assembly, where the plate comprises up to 96 wells.

10 The above assembly, where the plate comprises greater than 96 wells.

The above assembly, where the plate comprises greater than 384 wells.

The above assembly, where the electrodes are made of a material selected from the group consisting of gold, platinum, palladium, chromium, molybdenum, iridium, tungsten, tantalum and titanium.

15 The above assembly, where the electrodes are separated by a gap within the range of about 1 to 4 mm.

The above assembly, where the electrodes are separated by a gap within the range of about 0.1 to 1 mm.

20 The above assembly, where the electrodes are separated by a gap within the range of about 0.01 to 0.1 mm.

A bottom panel for a multi-well plate comprising.

at least one row of high transmittance regions with positions corresponding to well locations; a first: strip of conductive material extending along said row and overlapping a first portion of said well locations; and a second strip of conductive material extending along said row and overlapping a second portion of said well locations.

The above bottom panel, additionally comprising a first: electrical contact proximate to an end of said first strip and a second electrical contact proximate to an end of said second strip.

30 An assay apparatus comprising.

a sample well; a first pair of electrodes positioned within said sample well; at least one additional satellite electrode positioned within said sample well.

The above assay apparatus, where the at least one additional satellite electrode comprises second and third pairs of electrodes.

5 The above assay apparatus, where the satellite electrodes are charged to a potential less than that of the first pair of electrodes.

The above assay apparatus, where the electrodes are positioned with respect to the bottom surface of the well to provide an electric field adjacent to said bottom surface that varies by less than about 10% from a mean field intensity over at least about 20%
10 of the surface area of said bottom surface.

15 The present invention is not to be limited in scope by the specific embodiments described herein. Indeed, various modifications of the invention in addition to those described herein will become apparent to those skilled in the art from the foregoing description. Such modifications are intended to fall within the scope of the appended
20 claims.

Various publications are cited herein, the disclosures of which are incorporated by reference in their entireties. Furthermore, for general information, PCT Publication No. PCT/US01/21652 is incorporated herein in its entirety to the extent it is accurate and not inconsistent with the teachings herein. All patents, patent
25 applications, publications, texts and references discussed or cited herein are understood to be incorporated by reference to the same extent as if each individual publication or patent application was specifically and individually set forth in its entirety. In addition, all references, patents, applications, and other documents cited in an Invention Disclosure Statement, Examiner's Summary of Cited References, or
30 otherwise entered into the file history of this application are taken to be incorporated by reference into this specification for the benefit of later applications claiming

priority to this application. Finally, all terms not specifically defined are first taken to have the meaning given through usage in this disclosure, and if no such meaning is inferable, their normal meaning.

WHAT IS CLAIMED IS:

1. A method for identifying modulators of the activity of a voltage-gated ion channel comprising:
 - 5 (a) altering the transmembrane potential of at least a portion of the membrane of a cell expressing the voltage-gated ion channel by applying a voltage to the cells through extracellular electrodes while monitoring ion flow through the voltage-gated ion channel;
 - (b) exposing the cell in step (a) to a substance and monitoring ion
10 flow through the voltage-gated ion channel;
 - (c) comparing the ion flow through the voltage-gated ion channel in step (a) and step (b);
where a difference in the ion flow through the voltage-gated ion channel in step (a) and step (b) indicates that the substance is a modulator of the
15 voltage-gated ion channel.
2. A method for identifying modulators of the activity of a voltage-gated ion channel comprising:
 - 20 (a) dividing a plurality of cells expressing the voltage-gated ion channel into a control portion and a test portion;
 - (b) altering the transmembrane potential of the control portion of cells by applying a voltage to the cells through extracellular electrodes while monitoring ion flow through the voltage-gated ion channel;
 - (c) altering the transmembrane potential of the test portion of cells
25 by applying the voltage to the cells through extracellular electrodes in the presence of a substance while monitoring ion flow through the voltage-gated ion channel;
 - (d) comparing the ion flow through the voltage-gated ion channel in step (b) and step (c);
where a difference in the ion flow through the voltage-gated ion
30 channel in step (b) and step (c) indicates that the substance is modulator of the voltage-gated ion channel.
3. A method of identifying activators of a voltage-gated ion channel comprising:

- (a) providing a substrate upon which are living eukaryotic cells that express a plurality of the voltage-gated ion channels in their plasma membranes;
- (b) providing positive and negative electrodes positioned either on or near the substrate such that when a preselected voltage is applied through the positive and negative electrodes the transmembrane electrical potential of the cells is altered such that at least a portion of the voltage-gated ion channels are closed;
- (c) applying the preselected voltage through the positive and negative electrodes;
- (d) determining a control value for the flow of ions through the voltage-gated ion channels of the cells in step (c);
- (e) exposing the cells of step (c) to a substance for a period sufficient and under conditions such that a detectable number of the portion of the voltage-gated ion channels that are closed become open and allow ion flow through the detectable number of voltage-gated ion channels if the substance is an activator of the voltage-gated ion channels;
- (f) determining a test value for the flow of ions through the voltage-gated ion channels of the cells of step (e);
- (g) comparing the control value to the test value;
- where if the control value is less than the test value, then the substance is an activator of the voltage-gated ion channel.

4. The method of claim 3 where the substrate is glass or a multiwell tissue culture plate and is not silicon or a field effect transistor.
5. The method of claim 4 where the substrate contains wells in which the cells are present.
6. The method of claim 5 where the number of wells is 12, 24, 96, 384, 1,536, or 3,456.
7. The method of claim 5 where the wells are virtual wells.
8. The method of claim 3 where at least 50,000 substances are tested in a 24 hour period.

9. The method of claim 3 where the voltage-gated ion channel is a voltage-gated sodium channel, a voltage-gated potassium channel, or a voltage-gated calcium channel.

5

10. The method of claim 9 where the voltage-gated ion channel is a voltage-gated sodium channel.

11. The method of claim 9 where the voltage-gated ion channel is a voltage-gated potassium channel.

10

12. The method of claim 9 where the voltage-gated ion channel is a voltage-gated calcium channel.

15

13. The method of claim 3 where the cells are selected from the group consisting of: L cells L-M(TK⁻) (ATCC CCL 1.3), L cells L-M (ATCC CCL 1.2), HEK293 (ATCC CRL 1573), Raji (ATCC CCL 86), CV-1 (ATCC CCL 70), COS-1 (ATCC CRL 1650), COS-7 (ATCC CRL 1651), CHO-K1 (ATCC CCL 61), 3T3 (ATCC CCL 92), NIH/3T3 (ATCC CRL 1658), HeLa (ATCC CCL 2), C127I (ATCC CRL 1616), BS-C-1 (ATCC CCL 26), MRC-5 (ATCC CCL 171), CPAE (ATCC CCL 209), Saos-2 (ATCC HTB-85), ARPE-19 human retinal pigment epithelium (ATCC CRL-2302), GH3 cells, and primary cardiac myocytes.

20

14. The method of claim 13 where the cells are HEK293 (ATCC CRL 1573), GH3 cells, or primary cardiac myocytes.

25

15. The method of claim 3 where the cells contain a fluorescent indicator compound.

30

16. The method of claim 15 where the fluorescent indicator compound is selected from the group consisting of: fluo-3, fura-2, fluo-4, fluo-5, calcium green-1, Oregon green, 488 BAPTA, SNARF-1, and indo-1.

17. The method of claim 3 where the positive and negative electrodes are interdigitating.

18. The method of claim 3 where the substrate is a multiwell tissue culture plate having a plurality of wells that contain one positive and one negative electrode.

19. The method of claim 3 where the substrate is a multiwell tissue culture plate having a plurality of wells where one of the positive or negative electrodes forms the bottom of the wells and the other of the positive or negative electrode enters the wells from above.

20. The method of claim 3 where the substrate is a multiwell tissue culture plate having a plurality of virtual wells.

21. The method of claim 5 where each well contains from 10^3 to 10^7 cells and the cells contain a fluorescent indicator compound or a fluorescent voltage sensing dye.

22. The method of claim 3 where the cells do not naturally express the voltage-gated ion channel but have been transfected with an expression vector that encodes the voltage-gated ion channel.

23. A method of identifying inhibitors of a voltage-gated ion channel comprising:

- (a) providing a substrate upon which are living eukaryotic cells that express a plurality of the voltage-gated ion channels in their plasma membranes;
- (b) providing positive and negative electrodes positioned either on or near the substrate such that when a preselected voltage is applied through the positive and negative electrodes the transmembrane electrical potential of the cells is altered such that at least a portion of the voltage-gated ion channels are open;
- (c) applying the preselected voltage through the positive and negative electrodes;

- (d) determining a control value for the flow of ions through the voltage-gated ion channels of the cells in step (c);
- (e) exposing the cells of step (c) to a substance for a period sufficient and under conditions such that a detectable number of the portion of the voltage-gated ion channels that are open become closed and restrict ion flow through the detectable number of voltage-gated ion channels if the substance is an inhibitor of the voltage-gated ion channels;
- (f) determining a test value for the flow of ions through the voltage-gated ion channels of the cells of step (e);
- (g) comparing the control value to the test value;
where if the control value is greater than the test value, then the substance is an inhibitor of the voltage-gated ion channel.

24. The method of claim 23 where the substrate is glass or a multiwell tissue culture plate and is not silicon or a field effect transistor.

25. The method of claim 24 where the substrate contains wells in which the cells are present.

26. The method of claim 25 where the number of wells is 12, 24, 96, 384, 1,536, or 3,456.

27. The method of claim 26 where the wells are virtual wells.

28. The method of claim 23 where at least 50,000 substances are tested in a 24 hour period.

29. The method of claim 23 where the voltage-gated ion channel is a voltage-gated sodium channel, a voltage-gated potassium channel, or a voltage-gated calcium channel.

30. The method of claim 29 where the voltage-gated ion channel is a voltage-gated sodium channel.

31. The method of claim 29 where the voltage-gated ion channel is a voltage-gated potassium channel.

32. The method of claim 29 where the voltage-gated ion channel is
5 a voltage-gated calcium channel.

33. The method of claim 23 where the cells are selected from the group consisting of: L cells L-M(TK⁻) (ATCC CCL 1.3), L cells L-M (ATCC CCL 1.2), HEK293 (ATCC CRL 1573), Raji (ATCC CCL 86), CV-1 (ATCC CCL 70),
10 COS-1 (ATCC CRL 1650), COS-7 (ATCC CRL 1651), CHO-K1 (ATCC CCL 61), 3T3 (ATCC CCL 92), NIH/3T3 (ATCC CRL 1658), HeLa (ATCC CCL 2), C127I (ATCC CRL 1616), BS-C-1 (ATCC CCL 26), MRC-5 (ATCC CCL 171), CPAE (ATCC CCL 209), Saos-2 (ATCC HTB-85), ARPE-19 human retinal pigment epithelium (ATCC CRL-2302), GH3 cells, and primary cardiac myocytes.

15 34. The method of claim 33 where the cells are HEK293 (ATCC CRL 1573), GH3 cells, or primary cardiac myocytes.

35. The method of claim 23 where the cells contain a fluorescent
20 indicator compound.

36. The method of claim 35 where the fluorescent indicator compound is selected from the group consisting of: fluo-3, fura-2, fluo-4, fluo-5, calcium green-1, Oregon green, 488 BAPTA, SNARF-1, and indo-1.

25 37. The method of claim 23 where the positive and negative electrodes are interdigitating.

38. The method of claim 23 where the substrate is a multiwell
30 tissue culture plate having a plurality of wells that contain one positive and one negative electrode.

39. The method of claim 23 where the substrate is a multiwell tissue culture plate having a plurality of wells where one of the positive or negative

electrodes forms the bottom of the wells and the other of the positive or negative electrodes enters the wells from above.

40. The method of claim 23 where the substrate is a multiwell
5 tissue culture plate having a plurality of virtual wells.

41. The method of claim 25 where each well contains from 10^3 to
10 10^7 cells and the cells contain a fluorescent indicator compound or a fluorescent
voltage sensing dye.

42. The method of claim 23 where the cells do not naturally
express the voltage-gated ion channel but have been transfected with an expression
vector that encodes the voltage-gated ion channel.

43. A method of identifying activators of a voltage-gated ion
15 channel comprising:

- (a) providing a substrate upon which are living eukaryotic cells
that express a plurality of the voltage-gated ion channels in their plasma membranes;
- (b) providing positive and negative electrodes positioned either on
20 or near the substrate such that when a preselected voltage is applied through the
positive and negative electrodes the transmembrane electrical potential of the cells is
altered such that at least a portion of the voltage-gated ion channels are closed;
- (c) applying the preselected voltage through the positive and
negative electrodes to a control sample of the cells;
- 25 (d) determining a control value for the flow of ions through the
voltage-gated ion channels of the control sample of the cells in step (c);
- (e) applying the preselected voltage through the positive and
negative electrodes to a test sample of the cells while exposing the test sample of the
cells to a substance for a period sufficient and under conditions such that a detectable
30 number of the portion of the voltage-gated ion channels that are closed in the test
sample become open and allow ion flow through the detectable number of voltage-
gated ion channels if the substance is an activator of the voltage-gated ion channels;
- (f) determining a test value for the flow of ions through the
voltage-gated ion channels of the test sample of cells of step (e);

(g) comparing the control value to the test value;
where if the control value is less than the test value, then the substance is an activator of the voltage-gated ion channel.

- 5 44. A method of identifying inhibitors of a voltage-gated ion channel comprising:
- (a) providing a substrate upon which are living eukaryotic cells that express a plurality of the voltage-gated ion channels in their plasma membranes;
 - (b) providing positive and negative electrodes positioned either on
10 or near the substrate such that when a preselected voltage is applied through the positive and negative electrodes the transmembrane electrical potential of the cells is altered such that at least a portion of the voltage-gated ion channels are open;
 - (c) applying the preselected voltage through the positive and negative electrodes to a control sample of the cells;
 - 15 (d) determining a control value for the flow of ions through the voltage-gated ion channels of the control sample of the cells in step (c);
 - (e) applying the preselected voltage through the positive and negative electrodes to a test sample of the cells while exposing the test sample of the cells to a substance for a period sufficient and under conditions such that a detectable
20 number of the portion of the voltage-gated ion channels that are open in the test sample become closed and restrict ion flow through the detectable number of voltage-gated ion channels if the substance is an inhibitor of the voltage-gated ion channels;
 - (f) determining a test value for the flow of ions through the voltage-gated ion channels of the test sample of cells of step (e);
 - 25 (g) comparing the control value to the test value;
where if the control value is greater than the test value, then the substance is an inhibitor of the voltage-gated ion channel.

- 30 45. An apparatus for use in identifying activators or inhibitors of voltage-gated ion channels comprising:
a substrate having an upper surface upon which are present at least 10^3 living eukaryotic cells which have a voltage-gated ion channel of interest in their plasma membranes;

a plurality of positive electrodes and a plurality of negative electrodes positioned either on or near the substrate such that when a voltage is applied through the positive and negative electrodes the transmembrane potential of the cells is controlled;

- 5 at least one substance that is suspected of being an activator or an inhibitor of the voltage-gated ion channel;
 where the cells contain a fluorescent indicator compound.

46. A multiwell tissue culture plate having:
10 a plurality of wells in which are present at least 10^3 living eukaryotic cells per well of the plurality which cells have a voltage-gated ion channel of interest in their plasma membranes;

 a plurality of positive electrodes and a plurality of negative electrodes positioned such that when a preselected voltage is applied through the positive and
15 negative electrodes, the transmembrane potential of the cells is altered;

- at least one substance that is suspected of being an activator or an inhibitor of the voltage-gated ion channel in at least one of the plurality of the wells;
 where the cells contain a fluorescent indicator compound or a voltage sensitive membrane dye.

- 20 47. A multiwell tissue culture plate where a plurality of the wells of the plate contain a pair of electrodes disposed such that when a preselected voltage is applied across the electrodes the transmembrane potential of cells within the wells is altered.

- 25 48. The multiwell tissue culture plate of claim 47 where the multiwell tissue culture plate contains one of the pair of electrodes on the bottom of the wells and the other of the pair of electrodes on the side of the wells.

- 30 49. The multiwell tissue culture plate of claim 47 where the multiwell tissue culture plate contains both of the pair of electrodes on the bottom of the wells.

50. The multiwell tissue culture plate of claim 47 where one of the pair of electrodes is a layer of conductive material that forms the bottom of the wells and the other of the pair of electrodes enters the wells from above.

5 51. The multiwell tissue culture plate of claim 47 where both of the pair of electrodes are embedded in an insulator and enter the wells from above.

52. The multiwell tissue culture plate of claim 50 where the electrode that enters the wells from above has a central conductive material portion
10 that is surrounded by an insulator.

53. The multiwell tissue culture plate of claim 47 where the pairs of electrodes form an alternating pattern of positive and negative electrodes in the wells.
15

54. The multiwell tissue culture plate of claim 50 where the layer of conductive material that forms the bottom of the wells is a layer of indium tin oxide that overlays a glass substrate.

20 55. The multiwell tissue culture plate of claim 54 where the layer of conductive material and the glass substrate are transparent.

56. The multiwell tissue culture plate of claim 47 where a plurality of the wells of the plate contain interdigitating electrodes.
25

57. A multiwell tissue culture plate where:
the bottom of the wells is a filter membrane upon which cells can be grown;
the wells are located in a trough suitable for containing a fluid;
30 the trough contains a first electrode;
a second electrode enters the wells from above;
where the first and second electrodes are so disposed that when a preselected voltage is applied across the electrodes the transmembrane potential of cells within the wells is altered.

58. A combination of the multiwell tissue culture plate, as according to claims 46 to 57, and a fluorescence imager where the multiwell tissue culture plate and the fluorescence imager are positioned relative to one another such that the fluorescence imager can obtain fluorescence readings from the wells of the multiwell tissue culture plate.

59. A combination of a top substrate and a bottom substrate where the top and bottom substrates each contain:
a plurality of virtual wells; and
a layer of conductive material that forms the bottoms of the virtual wells;
where the layers of conductive material in the top and bottom substrates are connected to a pulse generator such that the layers of conductive material function as electrodes such that when a preselected voltage is applied across the electrodes the transmembrane potential of cells within the virtual wells is altered.

60. A substrate having square or rectangular wells formed by a plurality of generally parallel positive and negative electrodes and a plurality of spacers arranged generally at right angles to the electrodes, where:
one wall of the wells is formed by a positive electrode and the opposite wall of the well is formed by a negative electrode;
the spacers form the walls of the wells that are at right angles to the walls formed by the electrodes;
where the electrodes are so disposed that when a preselected voltage is applied across the electrodes the transmembrane potential of cells within the wells is altered.

61. A system for applying electrical field stimulation to cells, said system comprising:

a multiwell tissue culture plate, wherein the bottom of the wells is comprised of an *optically transparent* filter membrane upon which cells can be grown;
a trough suitable for containing fluid and configured such that said multiwell tissue culture plate may sit therein;

at least one first electrode disposed in said trough; and

an electrode head comprising a plurality of second electrodes in an amount
5 corresponding to the number of wells in said multiwell tissue culture plate, wherein
said electrode head and said plurality of said second electrodes are configured such
that said plurality of electrodes are disposed in the wells of the multiwell tissue
culture plate upon positioning said electrode head onto said multiwell tissue culture
plate;

10

wherein said at least one first electrode and said plurality of said second electrodes are
so disposed that when a preselected voltage is applied across the electrodes the
transmembrane potential of cells within the wells is altered.

15

62. The system of claim 61, further comprising a waveform generator
that is in electrical communication with said at least one first electrode or said
plurality of second electrodes, or both, whereby electric pulse signals are generated by
said waveform generator.

20

63. The system of claim 62 further comprising a computer electrically
connected to said waveform generator, said computer comprising software for
coordinating said pulse signals produced by said waveform generator.

25

64. The system of claim 62, wherein said waveform generator
generates a binary value that represents the address of the well to be excited by said
pulse signals.

30

65. The system of claim 62, further comprising electrical relays
upstream of said plurality of second electrodes.

66. The system of said 65 further comprising a microcontroller in
electrical communication with said waveform generator and said electrical relays, so
disposed such that upon receiving a trigger pulse and a particular binary value from
said waveform generator, said microcontroller switches on the appropriate relay

thereby directing a pulse to the particular electrode corresponding to said particular binary value.

67. The system of claim 61 wherein said trough comprises one first
5 electrode.

68. A system for applying electrical field stimulation to cells, said system comprising:

10 a multiwell tissue culture plate, wherein the bottom of the wells is comprised of a ^{transparent} filter membrane upon which cells can be grown;
optically

15 a tray comprising a plurality of individual troughs suitable for containing fluid;
wherein the number of said plurality of troughs corresponds to the amount of wells comprised in said multiwell tissue culture plate; wherein said plurality of troughs are so disposed to individually contain each well of said multiwell tissue culture plate; and wherein said plurality of troughs may be accessed by a port defined in said multiwell tissue culture plate and disposed laterally to each well;

20 a conductive electrode plate configured to be mounted above said multiwell tissue culture plate; wherein said electrode plate comprises a plurality of apertures configured to allow the wells of the multiwell tissue plate to pass through said conductive electrode plate; wherein said electrode plate comprises a plurality of
25 conductive pins integral or attached to said conductive electrode plate; and wherein individual pins of said plurality of conductive pins pass through said port to be disposed in individual troughs upon mounting said electrode plate on top of said multiwell tissue culture plate; and

30 an electrode head comprising a plurality of second electrodes in an amount corresponding to the number of wells in said multiwell tissue culture plate, wherein said electrode head and said plurality of said second electrodes are configured such that said plurality of electrodes are disposed in the wells of the multiwell tissue

culture plate upon positioning said electrode head onto said conductive electrode plate;

5 wherein said at least one first electrode and said plurality of said second electrodes are so disposed that when a preselected voltage is applied across the electrodes the transmembrane potential of cells within the wells is altered.

69. A novel electrode comprising a dielectric disc comprised of a dielectric material; a first conductive disc disposed on one side of said dielectric disc
10 and a second conductive disc disposed on the other side of said dielectric disc.

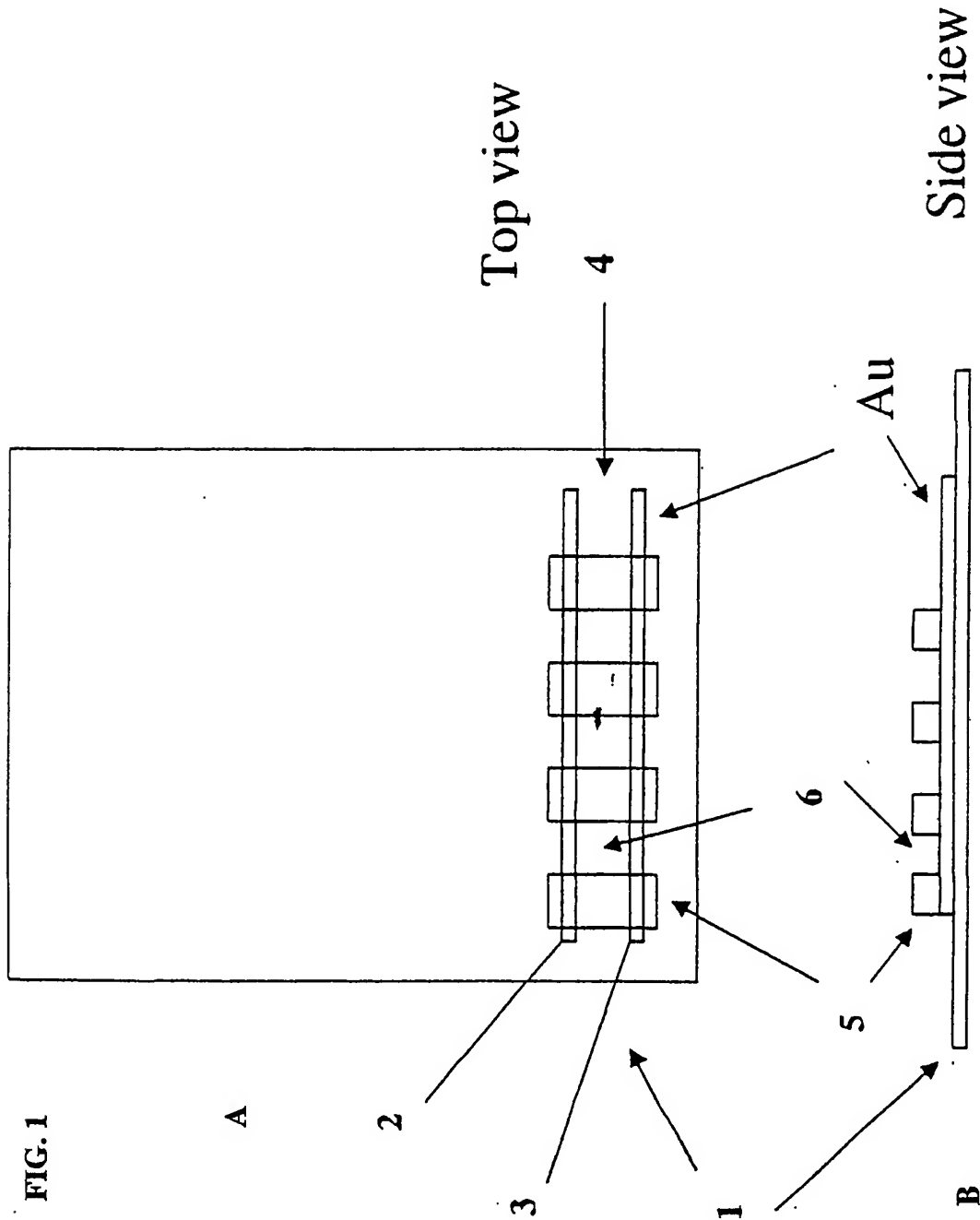
70. The electrode of claim 69 further comprising a concentric lead, wherein said concentric lead comprises at least one internal lead and at least one external lead whereby said internal lead passes through said first disc and said
15 dielectric disc and is electrically connected to said second disc.

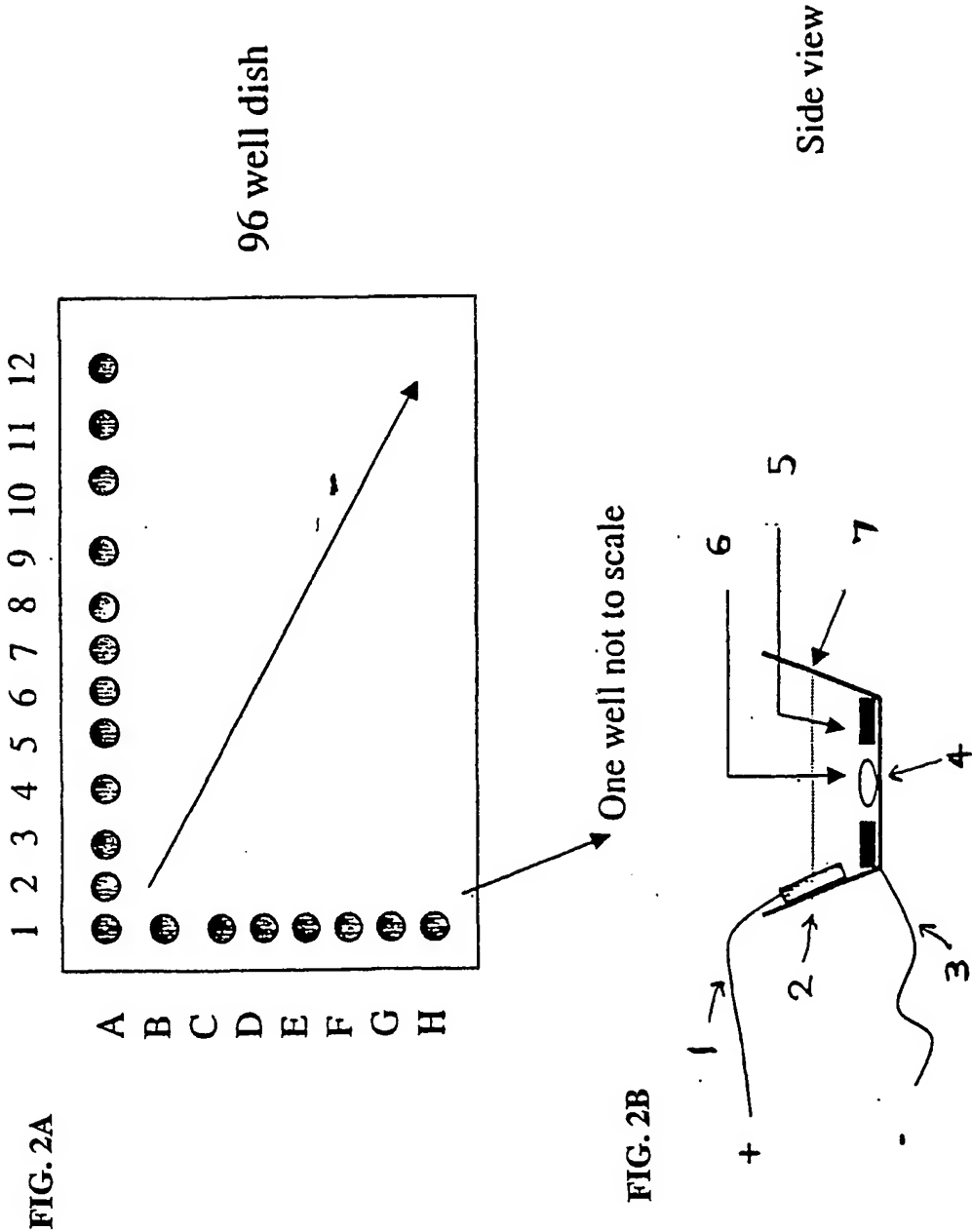
71. The electrode of claim 69 further comprising a first lead electrically connected to said first disc and a second lead electrically connected to said second disc.
20

72. The electrode of claim 69, wherein when a preselected voltage is applied across said first conductive disc and said second conductive disc to establish and electrical field.

25 73. The electrode of claim 72, wherein said electrode is able to provide a substantially uniform electrical field, while diminishing ohmic heating to a level such that said electrode may be brought into close proximity to cells to be manipulated.

30 74. The electrode of claim 73, wherein said electrode may be put in proximity with said cells at a distance of 10mm between said electrode and said cells to a distance closer to said cells without said electrode contacting said cells.





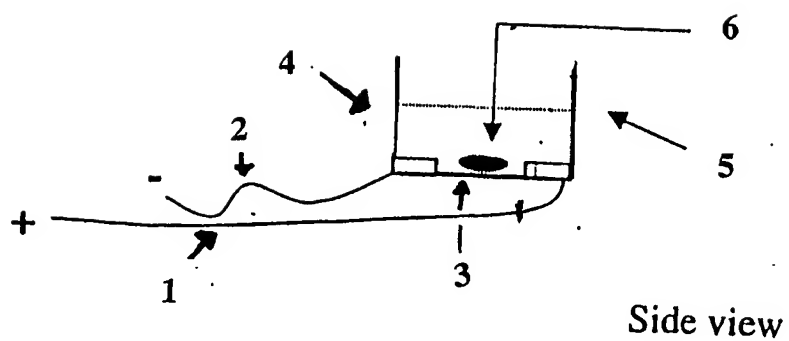


FIG. 2C

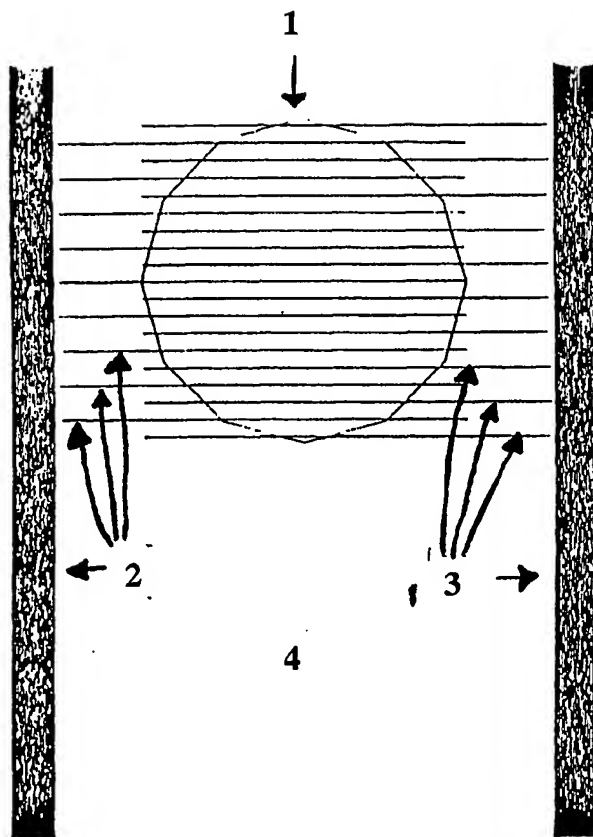


FIG. 3

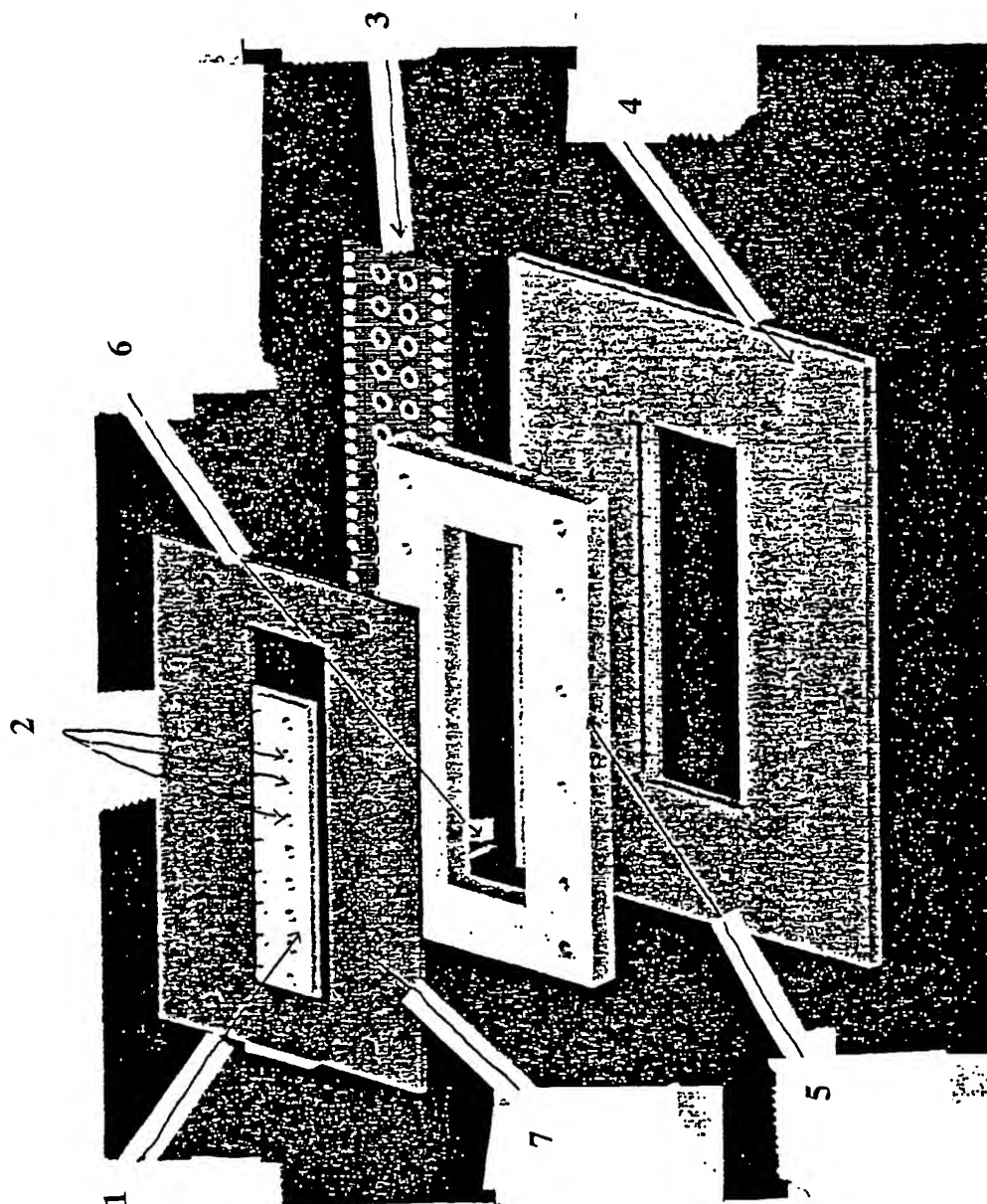


FIG. 4A

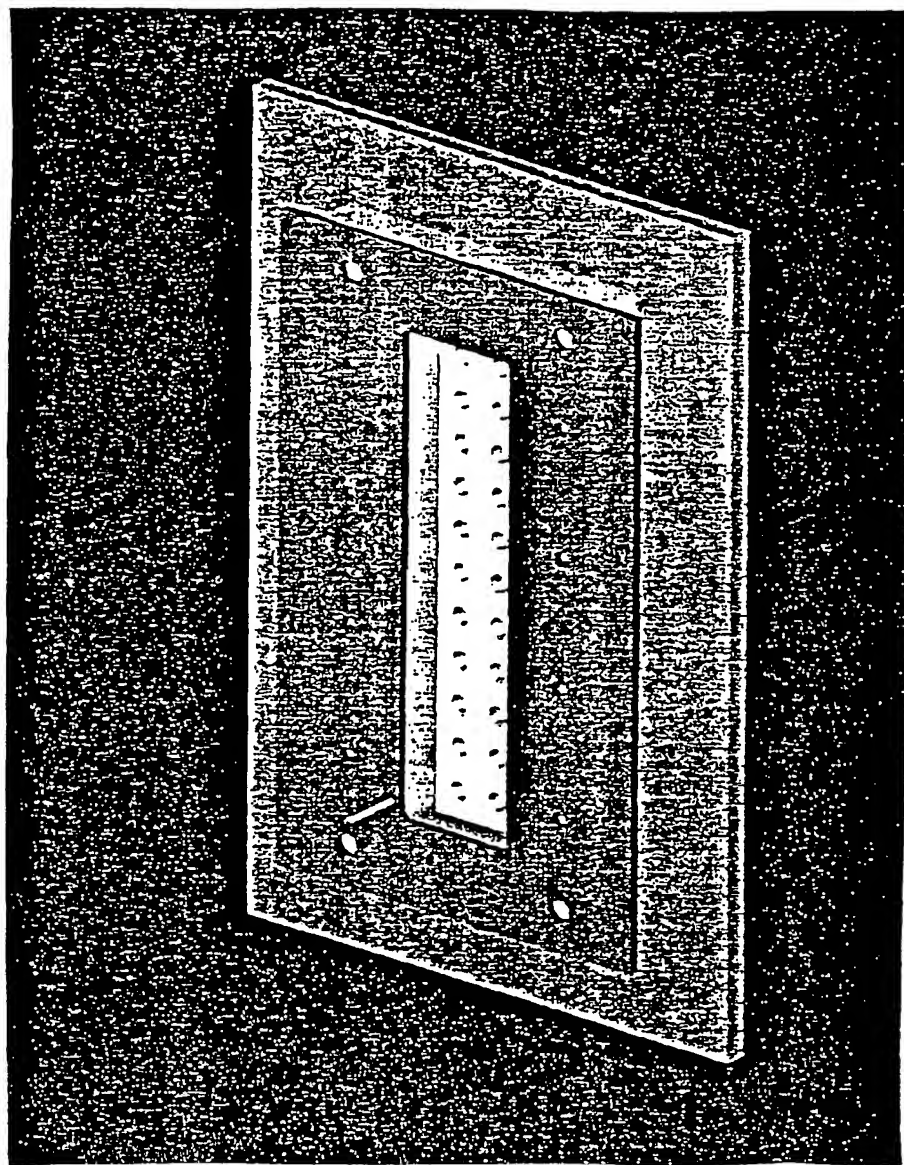


FIG. 4B

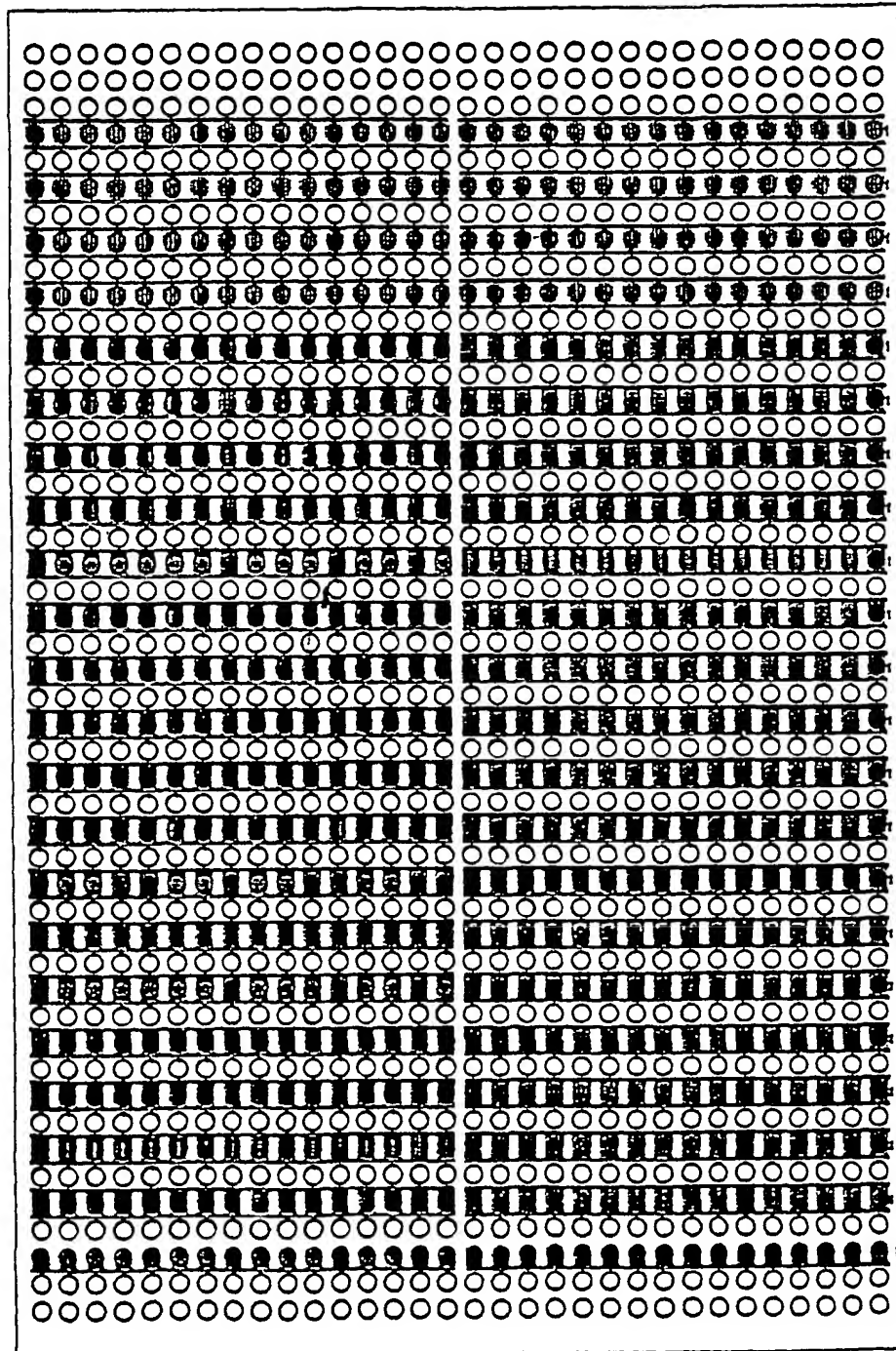


FIG. 5

8/56

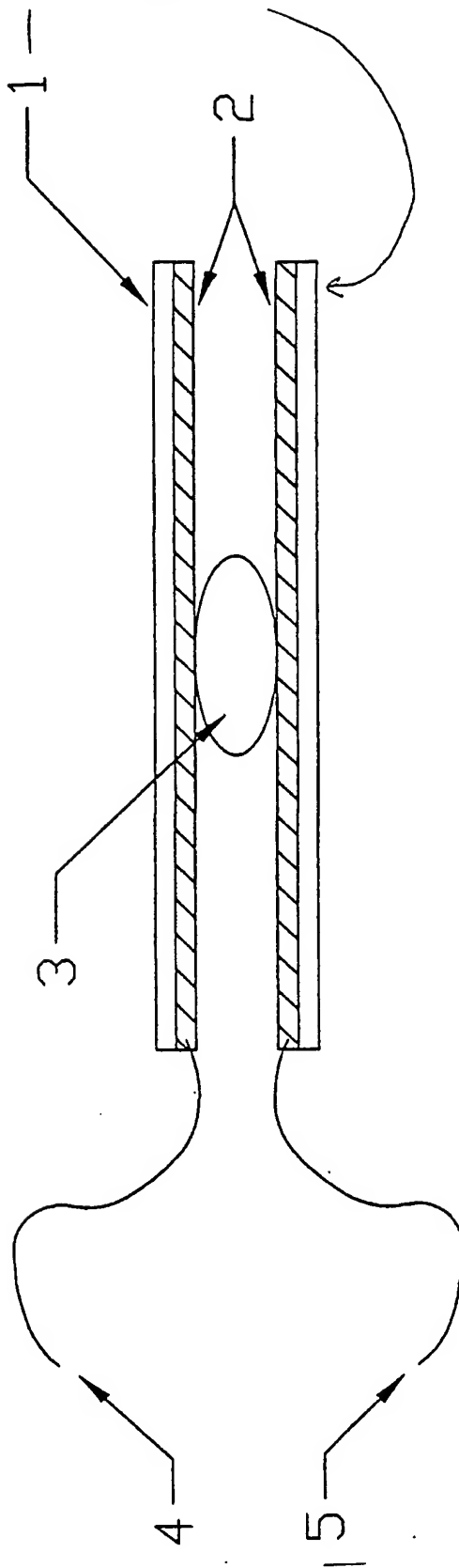


FIG. 6

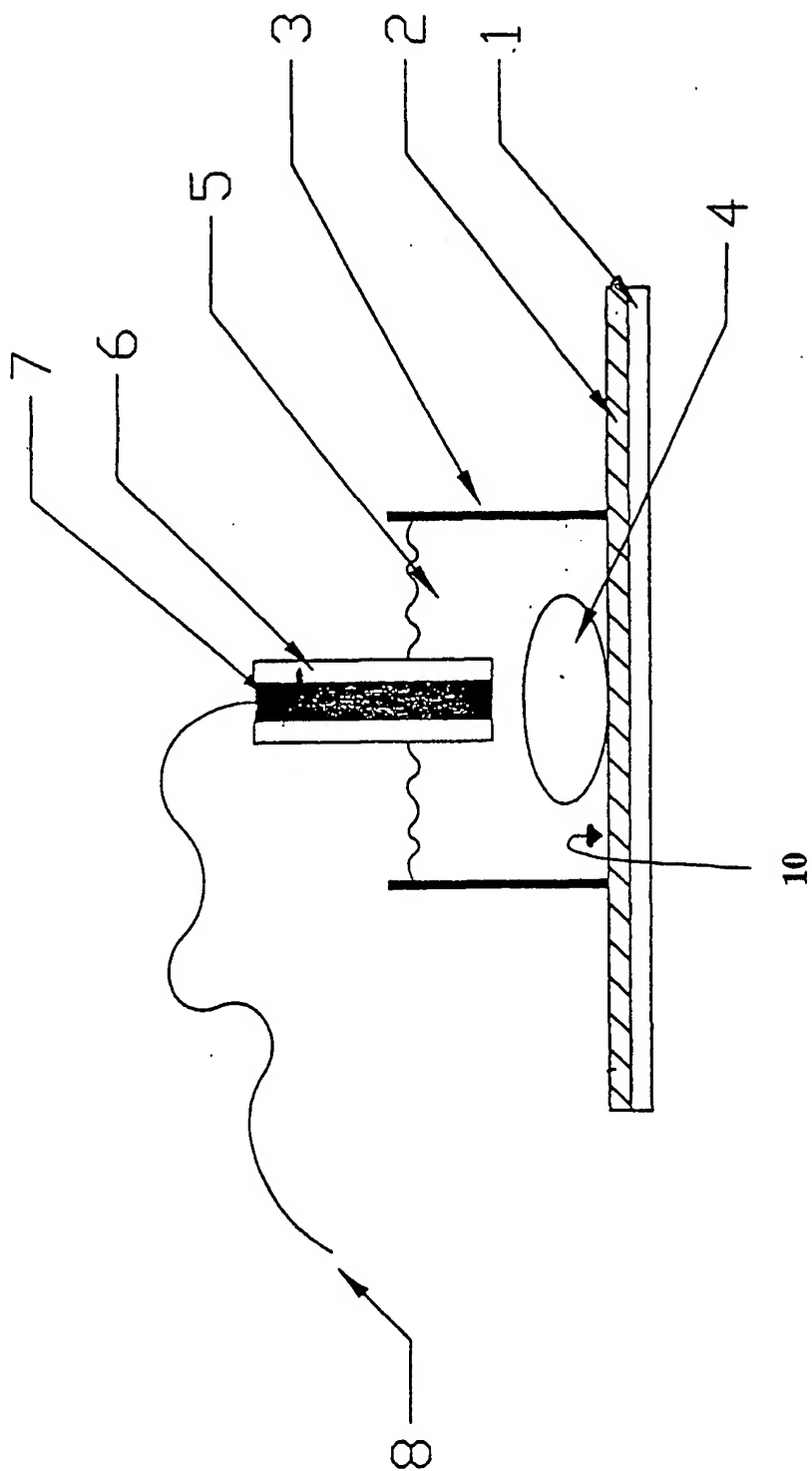


FIG. 7

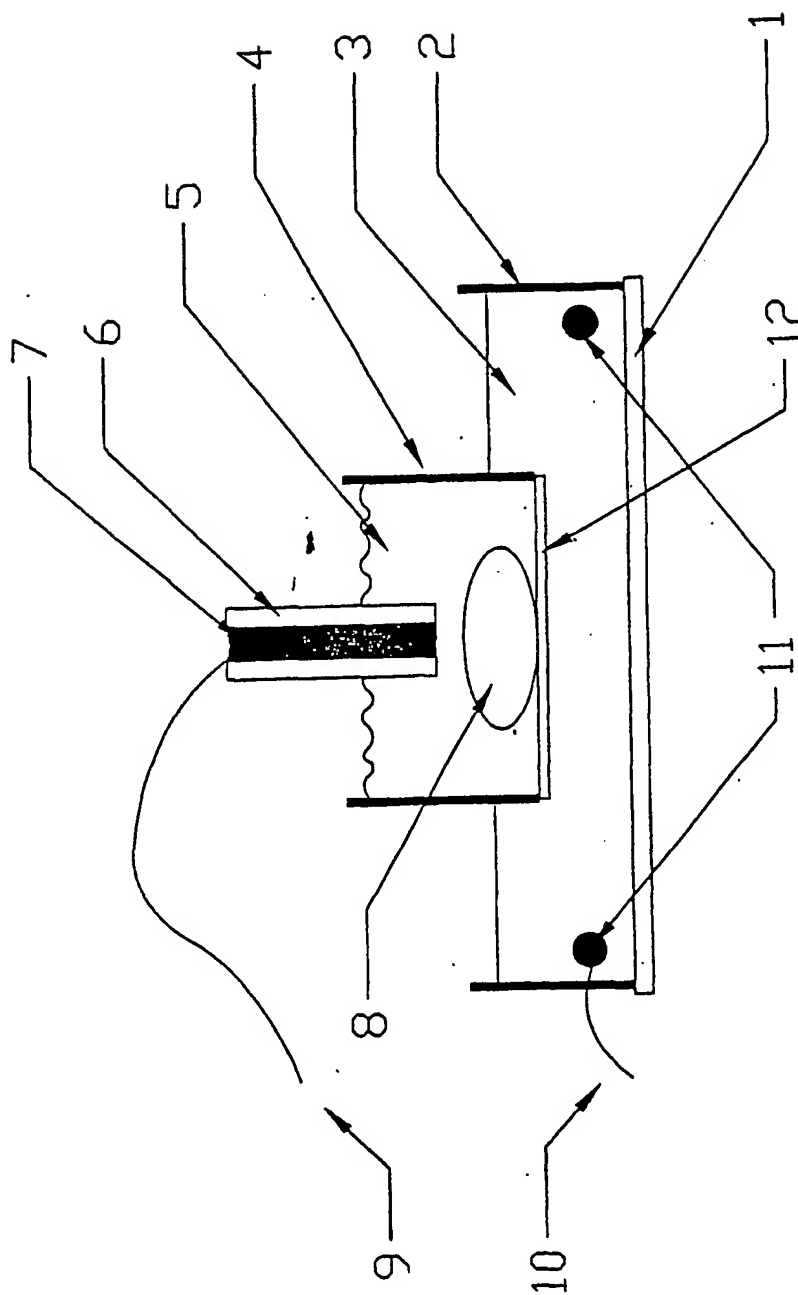
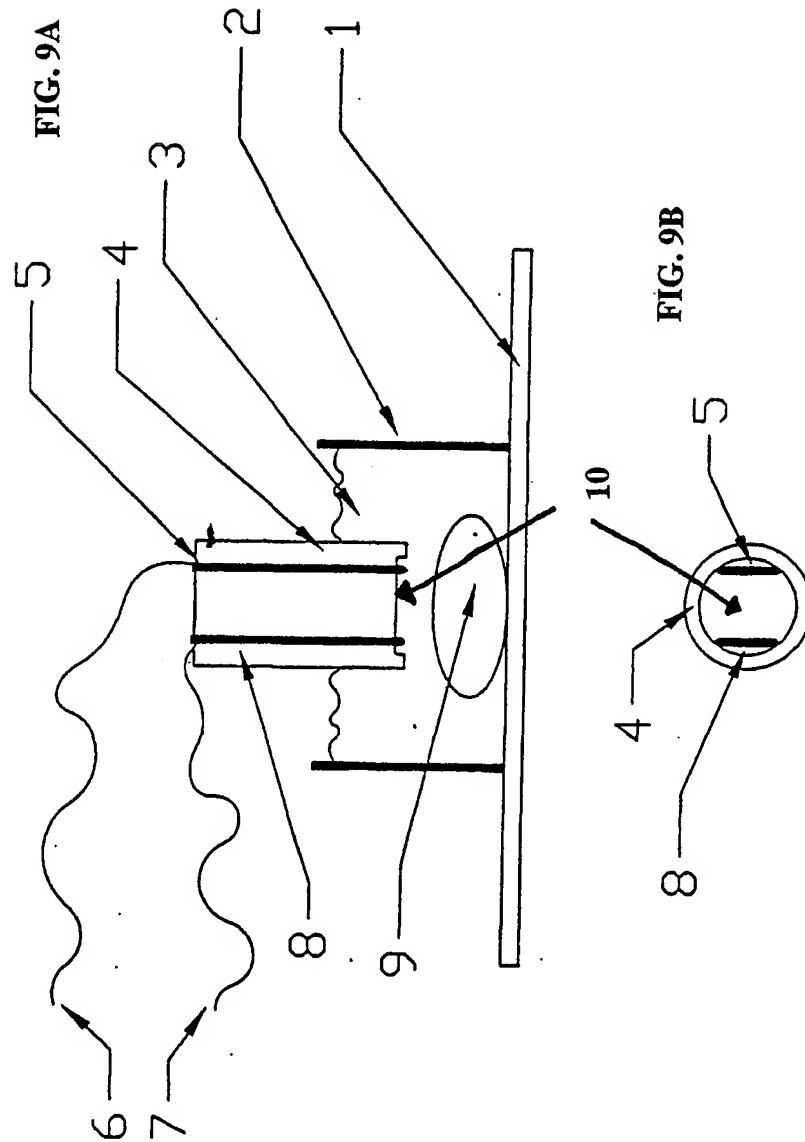
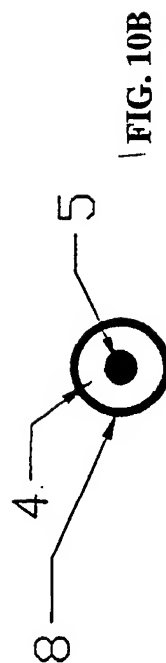
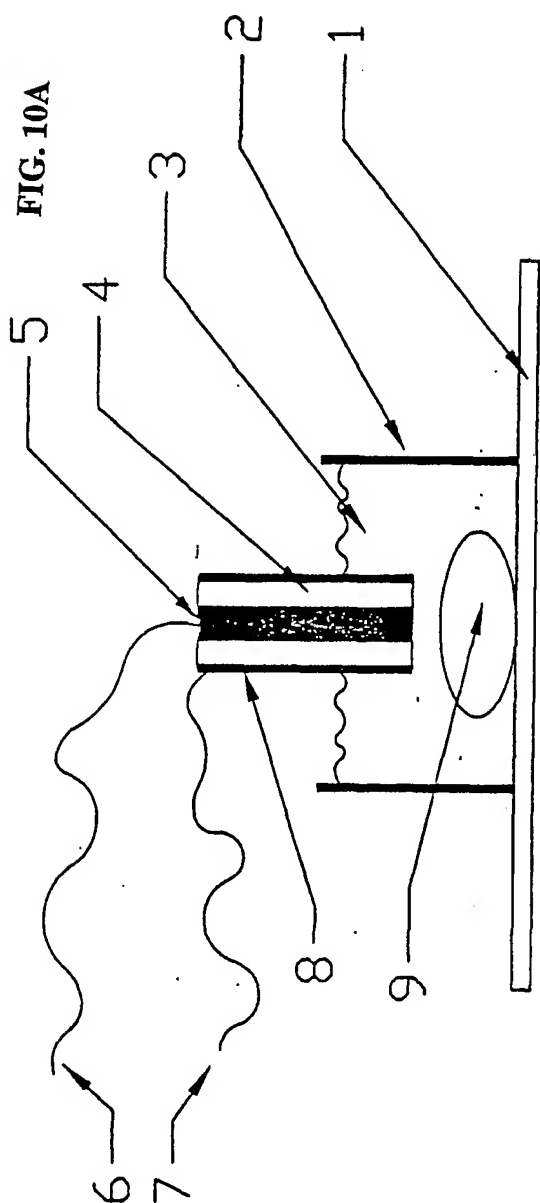


FIG. 8





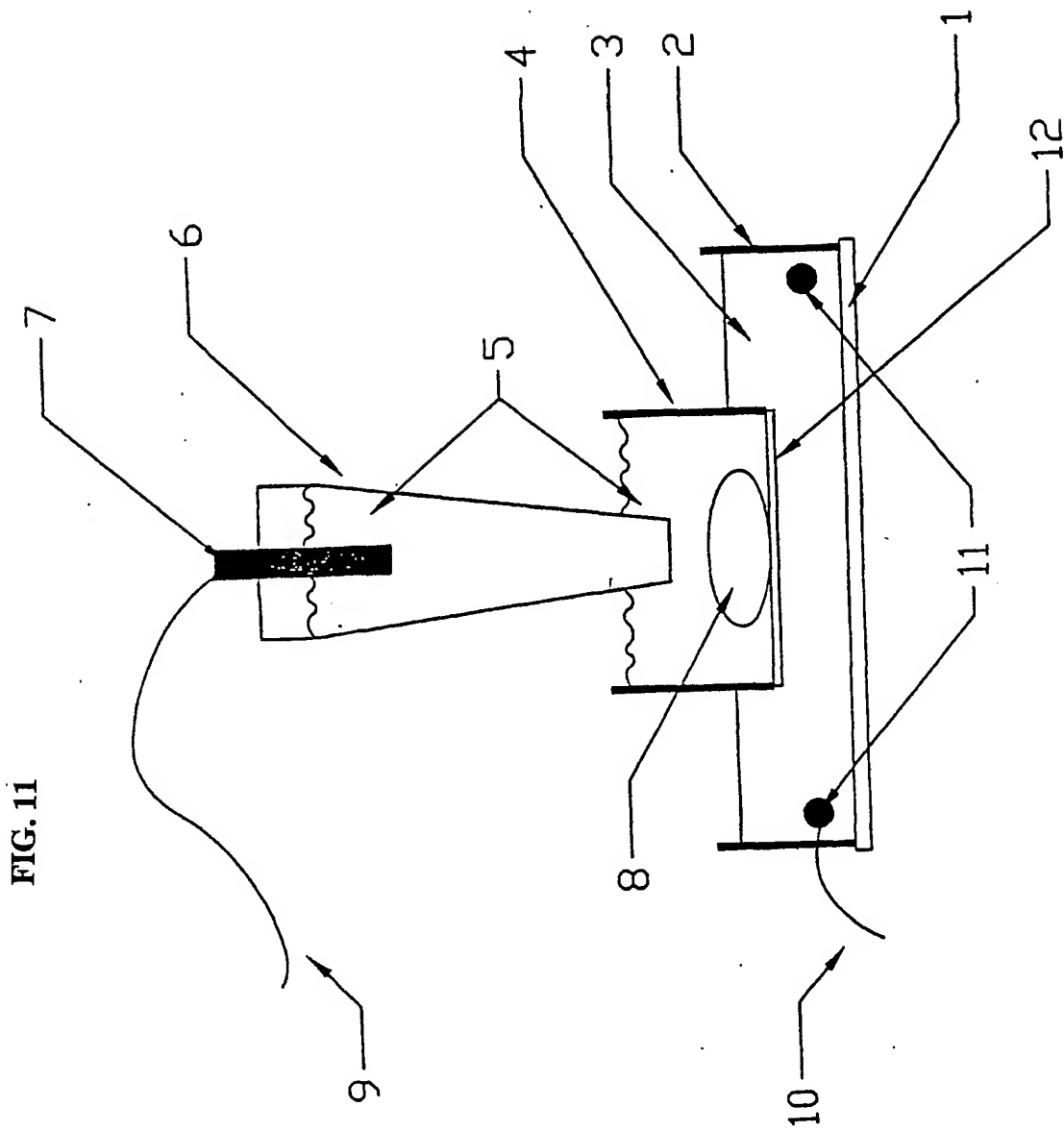


FIG. 11

FIG. 12A

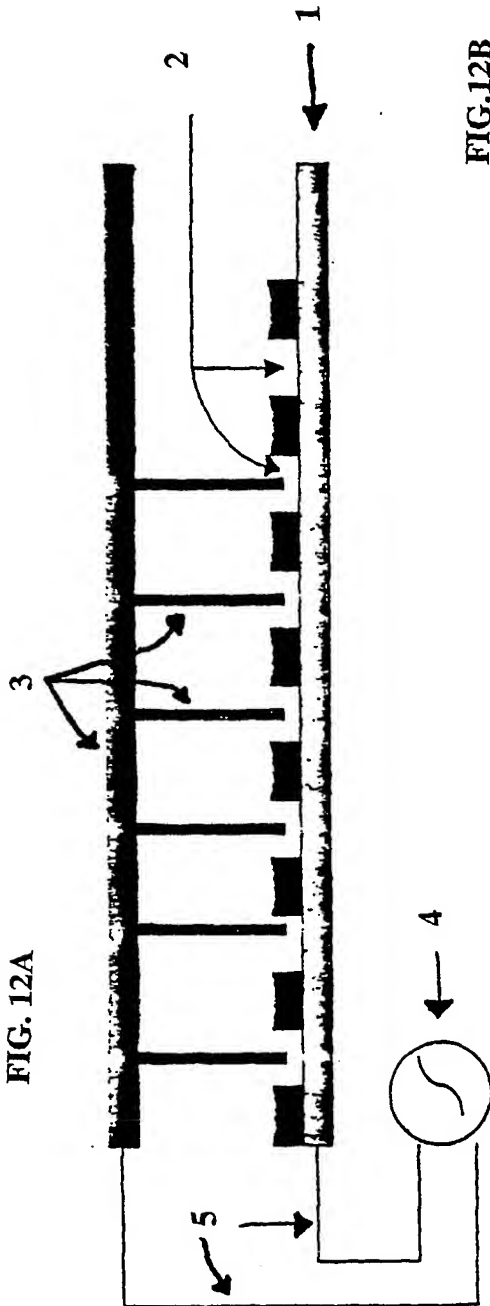


FIG. 12B

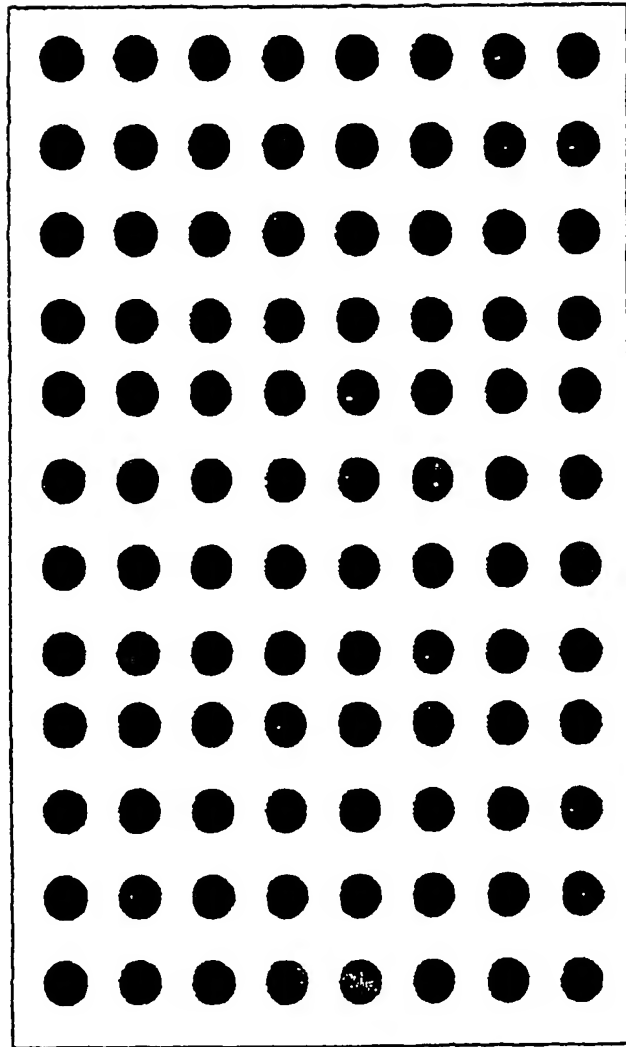


FIG. 13A

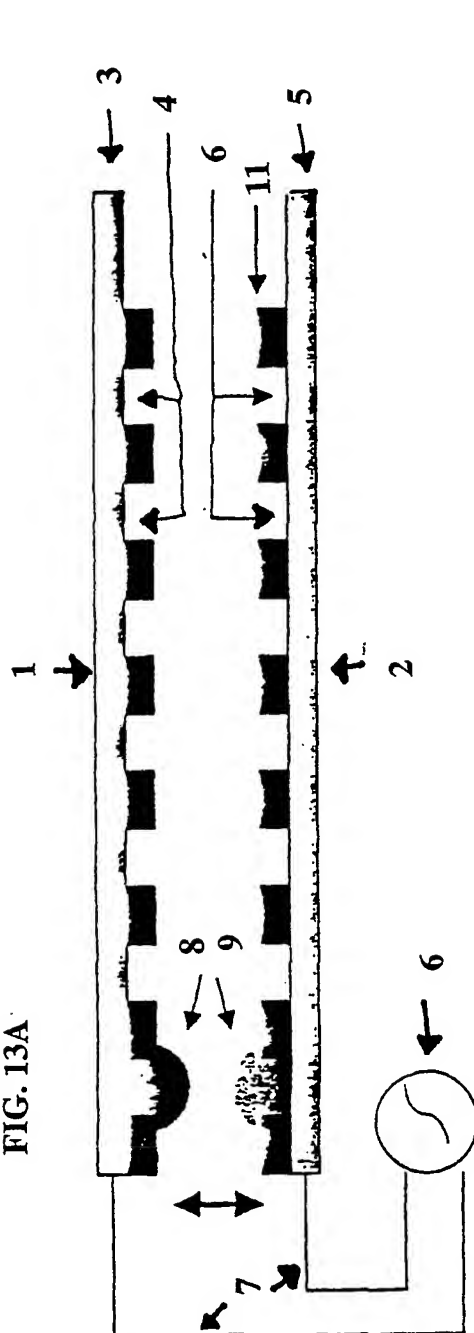
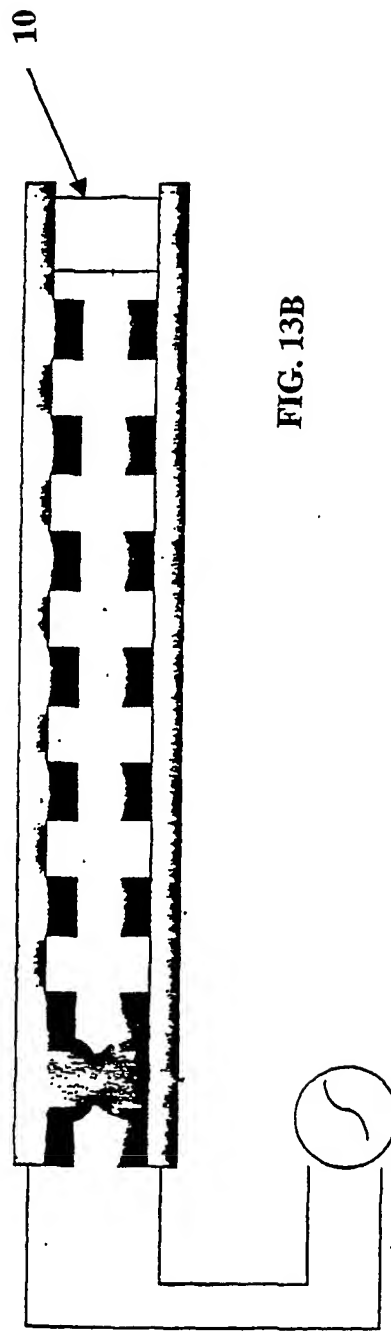


FIG. 13B



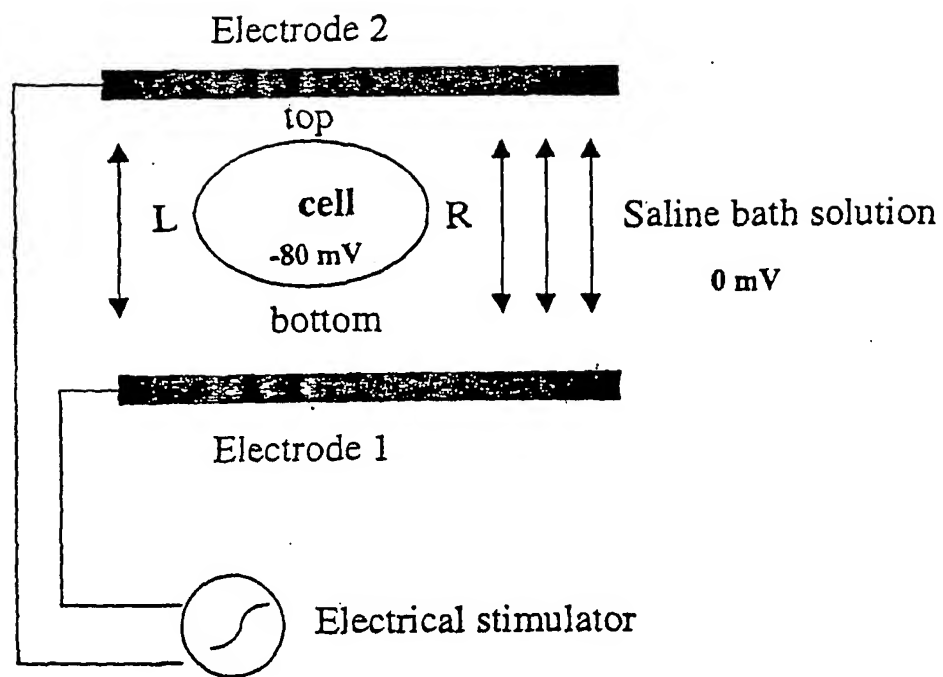


FIG. 14

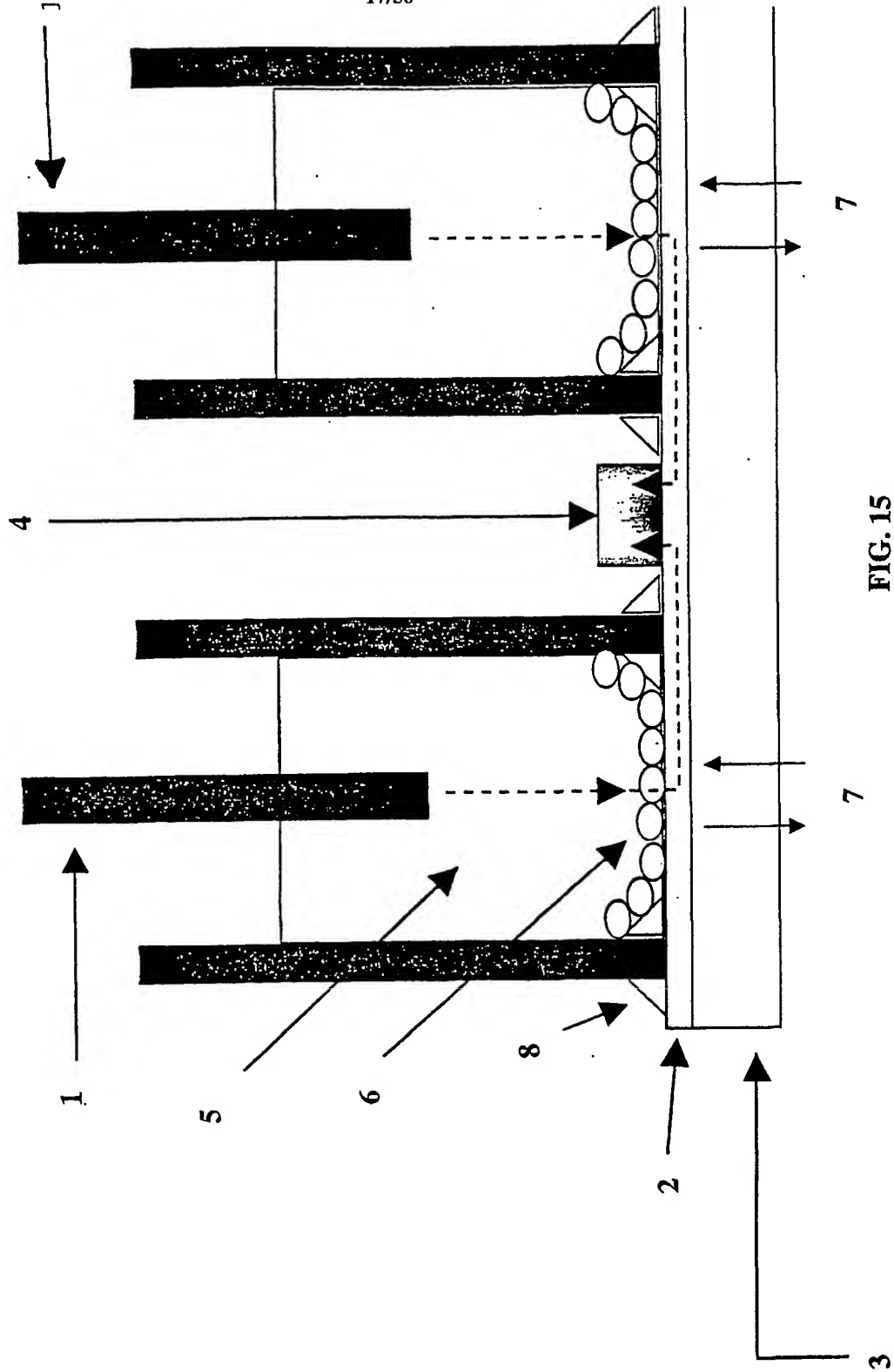
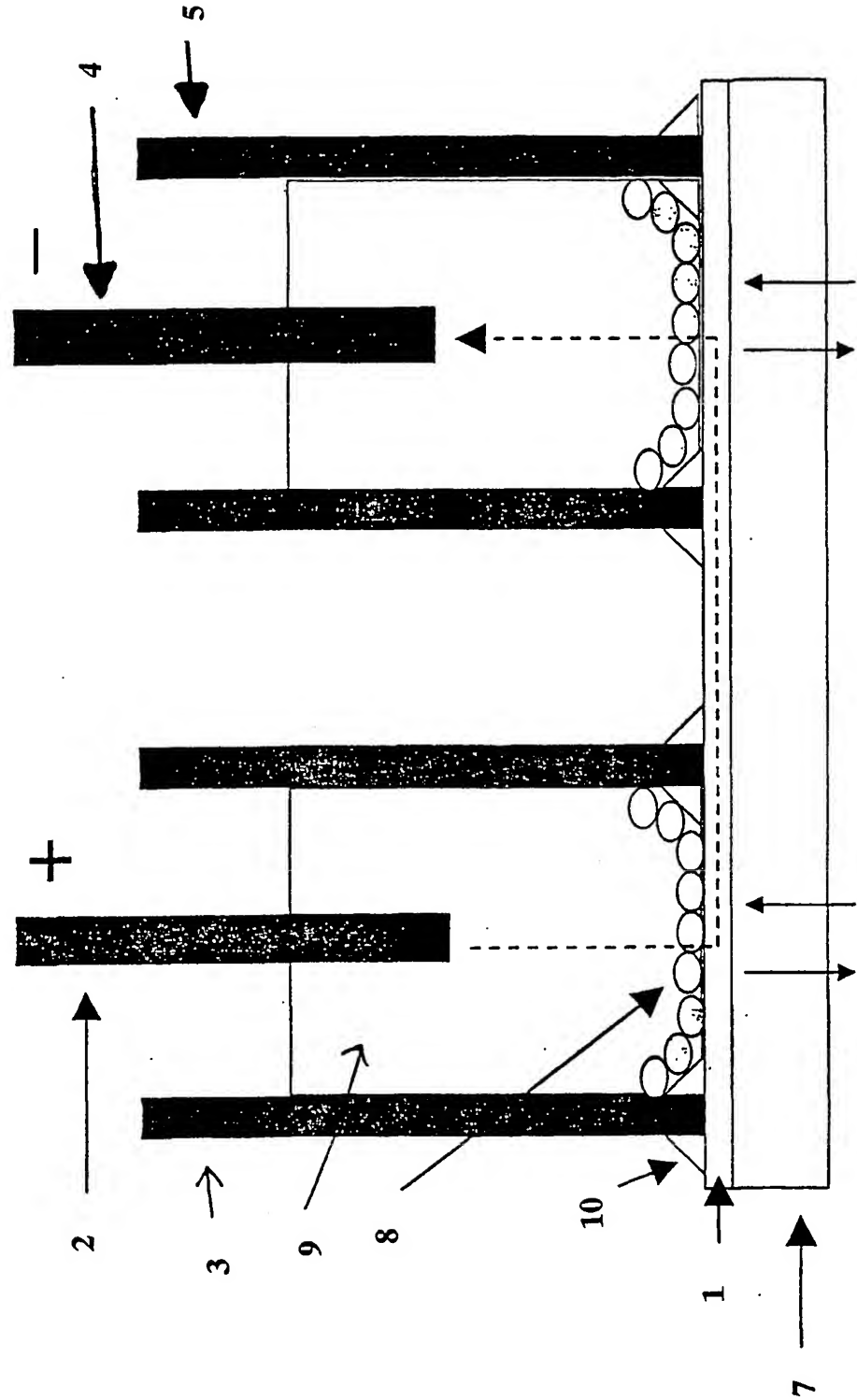


FIG. 16A



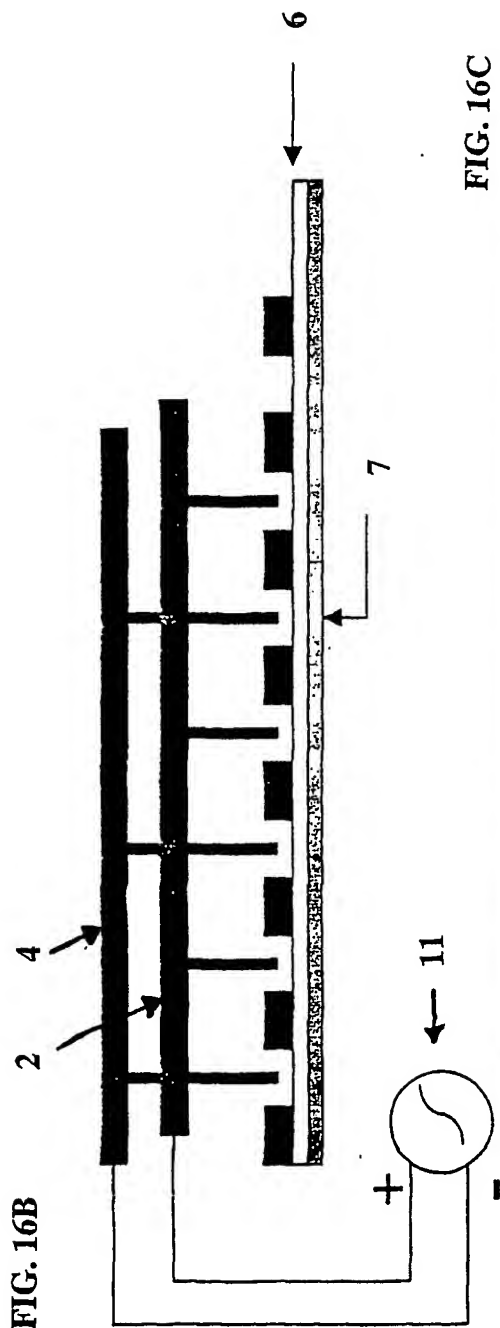


FIG. 16C

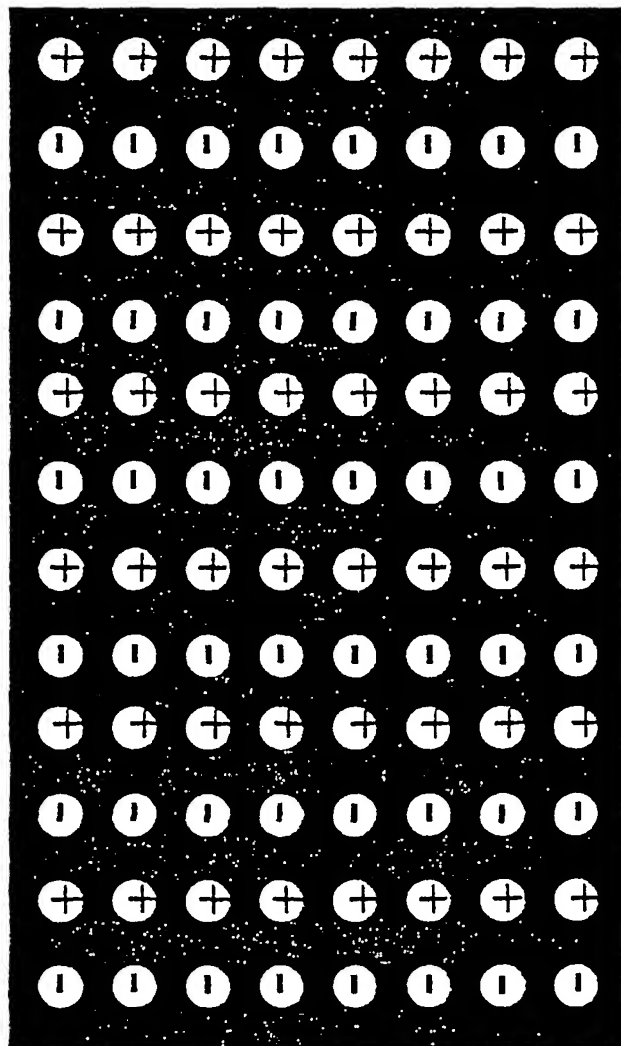


FIG. 16D

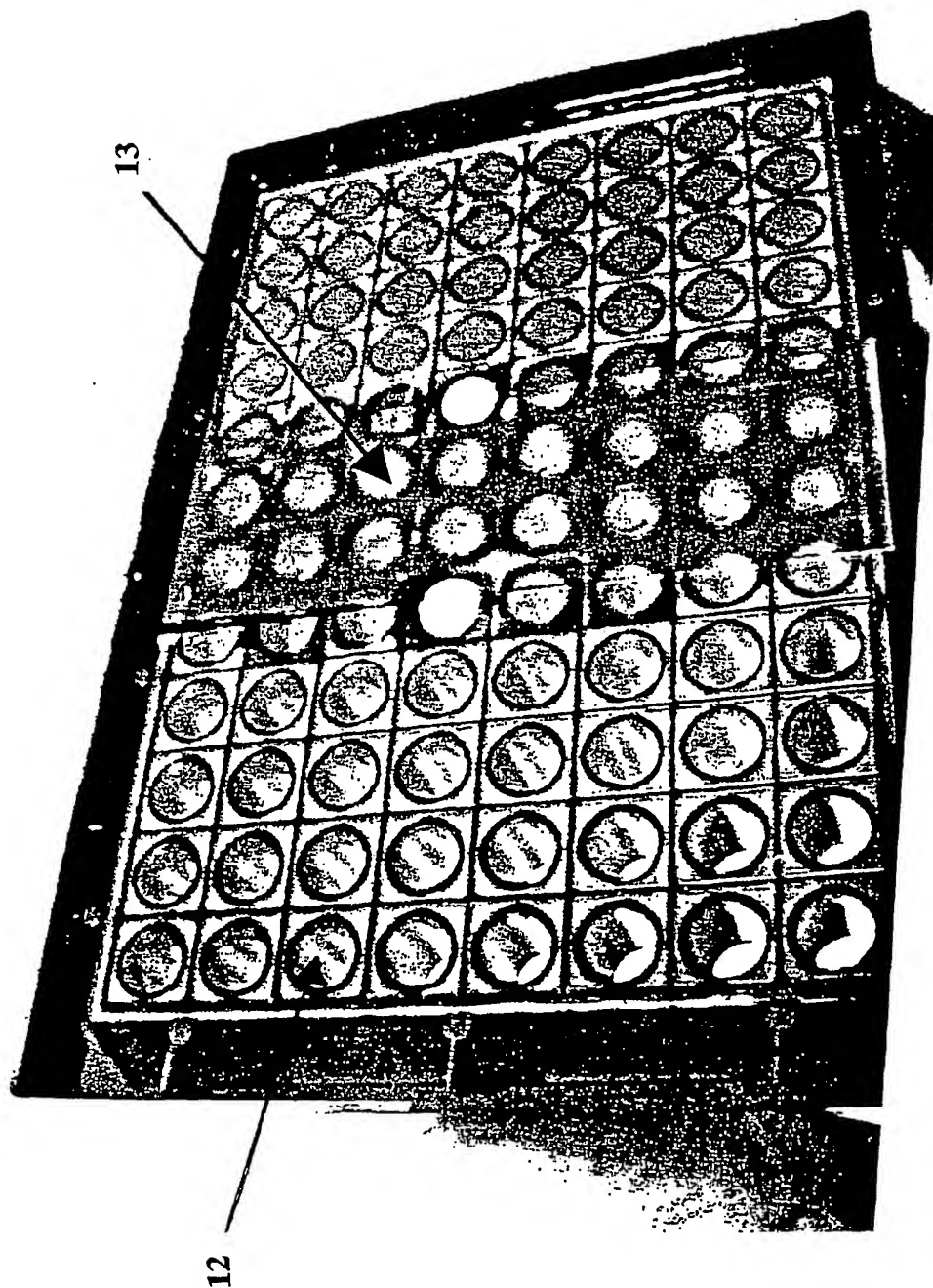


FIG. 17

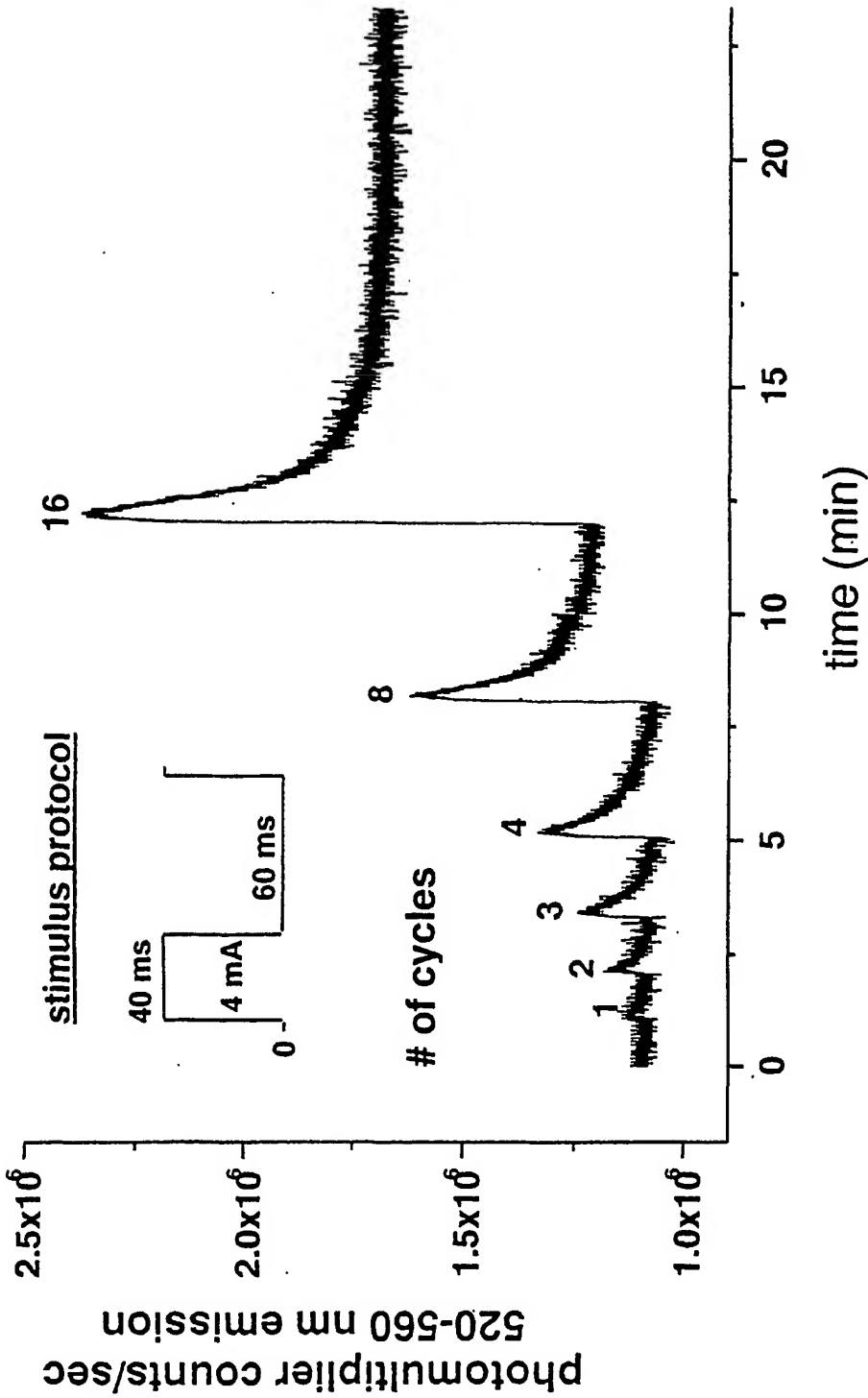


FIGURE 18A

1 atggaattcc ccattggatc cctcgaaact aacaacttcc gtcgctttac tccggagtca
 61 ctggtggaga tagagaagca aattgctgcc aagcagggaa caaagaaagc cagagagaag
 121 catagggagc agaaggacca agaagagaag cctcggcccc agctggactt gaaagcctgc
 181 aaccagctgc ccaagtcta tggtagctc ccagcagaac tgatcgggga gcccttggag
 241 gatctagatc cgttctacag cacacaccgg acatttatgg tgctgaaca agggaggacc
 301 atttcccgtt ttagtgccac tcgggccctg tggctattca gtcctttcaa cctgatcaga
 361 agaacggcca tcaaagtgtc tgtccactcg tggttcagtt tatttattac ggtcactatt
 421 ttggttaatt gtgtgtgcat gacccgaact gaccttccag agaaaattga atatgtctc
 481 actgtcattt acaccttga agccttgata aagatactgg caagaggatt ttgtctaaat
 541 gatttcacgt acctgagaga tccttggaaac tggctggatt ttagcgtcat taccctggca
 601 tatgttggca cagcaataga tctccgtggg atctcaggcc tgcggacatt cagagttctt
 661 agagcattaa aaacagtttc tgtgatccca ggccgtgaag tcattgtggg ggccctgatt
 721 cactcagtga agaaactggc tgatgtgacc atctcacca tcttctgctt aagtgtttt
 781 gccttgggtg ggctgcaact ctcaagggc aacctcaaaa ataaatgtgt caagaatgac
 841 atggctgtca atgagacaac caactactca tctcacagaa aaccagatat ctacataaat
 901 aagcgaggca cttctgaccc ctactgtgt ggcaatggat ctgactcagg ccactgccct
 961 gatggttata tctgccttaa aacttctgac aacccggatt ttaactacac cagctttgat
 1021 tcctttgctt gggttttctt ctactgttc gcctcatga cacaggattc ctgggaacgc
 1081 ctctaccagc agaccctgag gacttctggg aaaatctata tgatctttt tgtgctcgt
 1141 atcttcttgg gatcttctta cctggtaaac ttgatcttgg ctgtagtcac catggcgtat
 1201 gaggagcaga accaggcaac cactgatgaa attgaagcaa aggagaagaa gttccaggag
 1261 gccctcgaga tgctccggaa ggagcaggag gtgctagcag cactagggat tgacacaacc
 1321 tctctccact ccacaatgg atcaccttta acctcaaaa atgccagtga gagaaggcat
 1381 agaataaagc caagagtgtc agagggtcc acagaagaça acaaatcacc ccgctctgat
 1441 ccttacaacc agcgcaggat gtcttttcta ggctcgcct ctggaaaacg ccgggctagt
 1501 catggcagtg tgtccattt ccggtccct ggccgagata tctactccc tgaggagtc
 1561 acagatgatg gagtctttcc tggagaccac gaaagccatc ggggctctct gctgtgggt
 1621 gggggtgctg gccagcaagg cccctccctt agaagccctc ttctcaacc cagcaacctt
 1681 gactccaggc atggagaaga tgaacaccaa ccgcccacca ctagttagct tggccctgga
 1741 gctgtcgtat tctcggcatt cgatgcagga caaagaaga ctttctgtc agcagaatac
 1801 ttgatgaac cttccgggc ccaaagggca atgagtgtg tcagtatcat aacctccgtc
 1861 cttgaggaac tcgaggagtc tgaacagaag tgcccacctt gcttgaccag cttgtctcag
 1921 aagtatctga tctgggattg ctgccccatg tgggtgaagc tcaagacaat tcttttggg
 1981 cttgtgacgg atccctttgc agagctcacc atcaccttgt gcatcgttgt gaacaccatc
 2041 ttatggcca tggagacca tggcatgagc cctaccttgc aagccatgct ccagataggc
 2101 aacatcgtct ttaccatatt tttactgtt gaaatggctt tcaaatcat tgccttcgac
 2161 ccatactatt atttcagaa gaagtggaaat atctttgact gcatcatcgt cactgtgagt
 2221 ctgctagagc tggcggtggc caagaaggga agcctgtctg tgctcgggag cticcgttg
 2281 ctgcgcgtat tcaagctggc caaatccttg cccaccttaa acacactcat caagatcatc
 2341 ggaaactcag tgggggactt ggggaacctc accatcatcc tggccatcat tgtcttctc
 2401 ttgtctctgg ttggcaagca gctcctaggg gaaactacc gtaacaaccg aaaaaatc
 2461 tccgcgcccc atgaagactg gcccgttg ccatgcacg acttcttcca ctcttctc
 2521 attgtcttcc gtatccctg tggagagtgg attgagaaca tgtgggctg catggaagt
 2581 ggccaaaaat ccatacgct catcctttc ttgacggtga tgggtctagg gaacctgggt
 2641 gtgcttaacc tgttcatcgc cctgtattg aactcttca gtgctgaca cctcacagcc
 2701 ccggaggagc atggggaggt gaacaacctg caggtggccc tggcacggat ccaggtctt
 2761 ggccatcgt ccaaaccaggc tcttgcagc ttctcagca ggtcctgccc attccccag
 2821 cccaaggcag agcctgagct ggtggtgaaa ctccactct ccagctcaa ggctgagaac
 2881 cacattctc ccaacactc caagggagc tctggaggc tcaagctc cagaggcccc

FIG. 18B

2941 agggatgagc acagtgactt catcgctaata cgcactgtgt gggctctgt gcccattgct
 3001 gaggggaat ctgatcttga tgacttggag gatgatggg gggaagatgc tcagagcttc
 3061 cagcaggaag tgatcccaaa aggacagcag gagcagctgc agcaagtcca gaggtgtggg
 3121 gaccacctga caccaggag cccaggcact ggaacatctt ctgaggacct ggctccatcc
 3181 ctgggtgaga cgtggaaaaga tgagtctgtt cctcaggccc ctgctgaggg agtggacgac
 3241 acaagctcct ctgagggcag cacggtggac tgcctagatc ctgaggaaat cctgaggaag
 3301 atccctgagc tggcagatga cctggaagaa ccagatgact gcttcacaga aggatgcatt
 3361 cgccactgtc cctgtgcaa actggatacc accaagagtc catgggatgt gggctggcag
 3421 gtgcgcaaga cttgctaccg tatcgtggag cacagctggt ttgagagctt catcatcttc
 3481 atgatcctgc tcagcagtgg atctctggcc ttgaagactt attacctgga ccagaagccc
 3541 acggtgaaaag ctttgcctga gtacactgac agggctcttca cctttatctt tgtgttcgag
 3601 atgctgctta agtgggtggc ctatggcttc aaaaagtact tcaccaatgc ctggtgtctgg
 3661 ctggacttcc tcattgtgaa tatctactg ataagtcca cagcgaagat tctggaatat
 3721 tctgaagtgg ctcccatcaa agcccttcca acccttcgag cctgtcgccc actgcgggct
 3781 ctttctgat ttgaaggcat gcgggtgggt gtggatgccc tgggtggcgc catcccatcc
 3841 atcatgaatg tctctctgt ctgcctcatc ttctggctca tcttcagcat catgggtgtg
 3901 aacctctcg cagggaagt ttggaggtgc atcaactata ccgatggaga gttttccctt
 3961 gtacctttgt cgattgtgaa taacaagtct gactgcaaga ttcaaaactc cactggcagc
 4021 ttctctggg tcaatgtgaa agtcaacttt gataatgttg caatgggtta ccttgcaact
 4081 ctgcaggtgg caacctttaa aggtctggatg gacattatgt atgcagctgt tgattcccgg
 4141 gaggtcaaca tgcaacccaa gtgggaggac aacgtgtaca tgtatttgta ctttgcac
 4201 ttcatcattt ttggaggctt cttcacactg aatctcttg ttgggtcat aattgacaac
 4261 ttcaatcaac agaaaaaaaaa gttagggggc caggacatct tcatgacaga ggagcagaag
 4321 aaatactaca atgcatgaag gaagtgggc tccaagaagc cccagaagcc catcccagg
 4381 cccctgaaca agttccaggg tttgtctt gacatcgtga ccagacaagc tttgacatc
 4441 accatcatgg tctcatctg cctcaacatg atcccatga tgggtggagac tgatgacaa
 4501 agtgaagaaa agacgaaaat tctgggcaaa atcaaccagt tcttgtggc cgtcttcaca
 4561 ggcgaaatgt tcatgaagat gttcgtttg aggcagtact acttcacaaa tggctggaat
 4621 gtgttgactt tcattgtggt ggttctctcc attgcgagcc tgatttttc tgcaattctt
 4681 aagtcacttc aaagtactt cccccaaag ctttcagag tcatccgctt ggcccgaatt
 4741 ggccgcatcc tcagactgat ccgagcggcc aaggggatcc gcacactgct ctttgcctc
 4801 atgatgtccc tgcctgccc cttcaacatc gggctgttc tattccttgt catgttcac
 4861 tactccatct tcggtatgtc cagctttccc catgtgaggt gggaggctgg catcgacgac
 4921 atgttcaact tccagacctt cgccaacagc atgctgtgcc tcttcagat taccacgtc
 4981 gccggctggg atggcctct cagccccatc ctcaacacag ggcccccta ctgtgacccc
 5041 aatctgccc acagcaatgg caccagaggg gactgtggga gccagccgt aggcacatc
 5101 ttcttacc cctacatcat catctcctt ccatcgtgg tcaacatgta cattgcagt
 5161 attctggaga acttcaatgt ggccacggag gagagcactg agcctctgag tgaggacgac
 5221 ttgacatgt tcatgagac ctgggagaag ttgaccag aggcactca gtttattacc
 5281 ttttctgtc tctcgactt tgcagacat cttctgttc ccttgagaat cccaaaacc
 5341 aatcgaaata tactgatcca gatggacctg ctttggctc ctggagataa gatccactgc
 5401 ttgacatcc ttttgcctt caccaagaat gtcctaggag aatccgggga gttggattct
 5461 ctgaaggcaa atatggagga gaagttagt gcaactaat ttcaaaatc atcctatgaa
 5521 ccaatagcaa ccactctccg atggaagcaa gaagacatt cagccactgt cattcaaaag
 5581 gcctatcga gctatgtgt gcaccgtcc atggcactct ctaacacccc atgtgtgcc
 5641 agagctgagg aggaggctgc atactccca gatgaaggtt ttgtgcatt cacagcaat
 5701 caaaattctg tactcccaa caaatctgaa actcttctc ccacatcatt cccaccgtcc

FIG. 18C

MEFPIGSLETNNFRFTPESLVEIEKQIAAKQGTKKAREKHREQ
KDQEEKPRPQLDLKACNQLPKFYGELPAELIGELEDLDPFYSTHRTFMVLNKGRTIS
RFSATRALWLFSPFNLRRTAIKVSVHSWFSLFTTVTILVNCVCMTRTDLPEKIEYVF
TVIYTFEALIKILARGFCLNEFTYLRDPWNWLDIFSVTILAYVGTADLRGISGLRTFR
VLRALKTVSVIPGLKVTVGALHSVKKLADVLTILTFCLSVFALVGLQLFKGNLKNKC
VKNDMAVNETTNYSSHRKPDYINKRGTSDDLCCGNGSDSGHCPDGYICLKTSDNPFD
NYTSFDSFAWAFSLFRLMTQDSWERLYQQTLRTSGKIYMIFFVLVIFLGSFYLVNLI
LAVVTMAYEEQNQATTDEIEAKEKKFQEALEMLRKEQEVLAALGIDTTSLSHSHNGSPL
TSKNASERRHRIKPRVSEGSTEDNKSPRSDPYNQRRMSFLGLASGKRRASHGVSFHF
SPGRDISLPEGVTDDGVFPGDHESHRGSLLLGGGAGQQGPLPRSPLPQPSNPDSRHGE
DEHQPPPTSELAPGAVDVSADFAGQKKTFLSAEYLDEPFRAQRAMSVVSITSVLEEL
EESEQKCPPCLTSLSQKYLIWDCCPMWVKLKTILFGLVTDPAELTTTLCIVVNTIFM
AMEHHGMSPTFEAMLQIGNTVFTIFFTAEMVFKIIAFDPYFFQKKWNIFDCIIVTVS
LLELGVAKKGSLSVLRSFRLLRVFKLAKSWPTLNTLIKIGNSVGALGNLTILAITV
FVFALVGKQLLGENYRNNRKNISAPHEDWPRWHMHDFHSLIVFRILCGEWIENMWA
CMEVGQKSICLILFTVMVLGNLVVLNLFIALLLNSFSADNLTAPEDDGEVNNLQVAL
ARIQVFGHRTKQALCSFFSRSCPFPQPKAEPELVVKLPLSSSKAENHIAANTARGSSG
GLQAPRGPRDEHSDFIANPTVWVSVPIAEGESDLDLEDGDDGEDAQSFQQEVIPKGQQ
EQLQQVERCGDHLTPRSPGTGTSSD LAPSLGETWKDESVPQAPAEGVDDTSSEGST
VDCLDPEEILRKIPELADDLEEDDCFTGECIRHCPCKLDTTKSPWDVGWQVRKTCY
RIVEHSWFESFIIFMILLSSGSLAFEDYYLDQKPTVKALLEYTDRTVFTFIFVEMLLK
WVAYGFKKYFTNAWCWLDLIVNISLISLTAKILEYSEVAPIKALRTLRLRPLRALS
RFEGMRVVVDALVGAIPSIMNVLLVCLIFWLIFSIMGVNLFAKFWRCINYTDGEFSL
VPLSIVNNKSDCKIQNSTGSFFWVNVKVNFDNVAMGYLALLQVATFKGWMDMYAAVD
SREVMQPKWEDNVYMYLYFVIFIFGGFFTLNLFVGVIIDNFNQKKKLGGQDIFMT
EEQKKYYNAMKKLGSKKPQKPIRPLNKFQGFVFDIVTRQAFDITIMVLICLNMITMM
VETDDQSEKTKILGKINQFFVAVFTGECVMKMFALRQYYFTNGWNVFDFTVVVLSIA
SLIFSAILKSLQSYFSPTLFRVIRLARIGRILRLIRAAGKIRTLLFALMMSLPALFNI
GLLLFLVMFTYSIFGMSSFPVHVRWEAGIDDMFNFTQTFANSMCLCFQITTSAGWDGLLS
PILNTGPPYCDPNLPNSNGTRGDCGSPA VGIIFFTTYIISFLIVVNMYIAVILENFN
VATEESTEPLSEDDFDMFYETWEKFDPEATQFITFSALSDFADTSLGPLRIPKPNRNI
LIQMDLPLVPGDKIHCLDILFAFTKNVLGESGELDSLKANMEEKFMATNLSKSSYEPI
ATTLRWKQEDISATVIQKAYRSYVLHRSMALSNTPCVPRAEEEEASLPDEGFVAFTAN
ENCVLDPKSETASATSFPPSYESVTRGLSDRVNMRTSSSIQNEDEATSMELIAPGP

FIGURE 19A

1 cgaggccgcc gccgtgcct ccgccggcg agccggagcc ggagtcgagc cgcggccggg
61 agccgggagg gctggggacg cgggcccggg gcggaggcgc tggggggccgg ggccgggggg
121 gggggcggag gcgctggggg ccggggccgg gcccgggcgc cagcgggggt ccgcgtgac
181 cgcgccgcc gggcgatgcc cgcggggacg ccgccggcca gcagagcgag gtgtgccgg
241 ccgccacat gaccgagggc gcacggggcc ccgacgaggt ccgggtgcc ctgggcgcgc
301 cggcccttg cctgcggcg ttgtggggg cgtcccga gagccccgg gcgccgggac
361 gcgagcgga gcgggggtcc gagtcggcg tgcaccctc cgagagccc gcggccgagc
421 gcggcgcgga gctgggtgcc gacgaggagc agcgcgtccc gtaccggcc ttggcgcca
481 cggtcttct ctgcctcgg cagaccacgc ggccgcgag ctggtgcctc cggctggtc
541 gcaacccatg gttcagcac gtgagcatgc tggtaacat gctaacatgc gtgaccctg
601 gcatgttcg gccctgtgag gacgttgagt gcggctccga gcgctgcaac atcctggagg
661 ccttgacgc ctcatcttc gcctttttg cggtgagat ggtcatcaag atggtggcct
721 tgggctgtt cgggcagaag tttacctgg gtgacacgtg gaacaggctg gatttttca
781 tctcgtggc gggcatgatg gactactct tggacggaca caactgagc ctctcggta
841 tcaggaccgt gcgggtgctg cggccctcc gcgccatcaa ccgctgcct agcatgcgga
901 tcttggtcac tctgtgctg gatacgtgc ccatgctcg gaacgtcct ctgctgtgt
961 tctcgtctt ctcatcttc ggcatcgtg gcgtccagct ctgggtggc ctctgcgga
1021 accgtgctt cctggacagt gccttgtca ggaacaaca cctgacctc ctgcggcgt
1081 actaccagac ggaggagggc gaggagaacc cgtcatctg ctctcagc cgagacaacg
1141 gcatgcagaa gtgtcgcac atccccggc gccgcgagct gcgcatgcc tgcacctg
1201 gctgggaggc ctacacgcag ccgcaggcc aggggggtgg cgctgcacgc aacgcctgca
1261 tcaactgga ccagtactac aactgtgcc gtcgggtga ctcaacccc cacaacggtg
1321 ccatcaact cgacaacatc ggctacgct ggattgcat ctccagggtg atcacgtgg
1381 aaggctgggt ggacatcatg tactacgca tggacggcca ctactctac aactcatct
1441 attcatcct gctcatcatc gtgggtcct tctcatgat caactgtgc ctggtggtga
1501 ttgccacgca gttctcggag acgaagcagc ggagagagtc gctgatcgg gagcagcggg
1561 cagccacct gtccaacgac agcacgctgg ccagctctc cagcctggc agctgtacg
1621 aagagctgt gaagtacgtg ggccacatat tccgaaggt caagcggcgc agcttgcgc
1681 tctacggcg ctggcagagc cgctggcgca agaagggtga cccagtgct gtcaaggcc
1741 aggggtccgg gcaccggcag cggcgggcag gcaggcacac agcctcgtg caccacctg
1801 tctaccacca ccatcaccac caccaccac actaccatt cagccatggc agccccgca
1861 ggccggccc cgagccaggc gcctgcgaca ccaggctgtt ccgagctggc gcggccccct
1921 cgccacctc ccaggccgc ggacccccg acgagagtc tgtcacagc atctaccatg
1981 ccgactgcca catagagggg ccgcaggaga gggccgggt ggcacatgcc gcagccactg
2041 ccgctccag cctcaggctg gccacagggc tgggcacat gaactacccc acgacctgc
2101 cctcagggt gggcagcggc aaaggcagca ccagccccg acccaagggg aagtgggccc
2161 gtggaccgcc aggcaccggg gggcacggcc cgttgagctt gaacagccct gatccctacg
2221 agaagatccc gcatgtgttc ggggagcatg gactgggcca ggccctggc catctgtcg
2281 gcctcagtgt gccctgccc ctgcccagc cccagcggg cacttgacc tgtgagctga
2341 agagctgccc gtactgcacc cgtgcctgg aggacccgga ggtgagctc agcggtcgg
2401 aaagtggaga ctcatatggc cgtggcgtct atgaattac gcaggacgtc cggcacggtg
2461 accgtggga cccacgcga ccacccgtg gcagggacac accaggccca ggccaggca
2521 gccccagcg gcgggcacag cagaggggcag cccggggcga gccaggctgg atggggccc
2581 tctgggttac ctacagcgc aagctgcgc gcatcgtgga cagcaagtac ttcagccgtg
2641 gcatcatgat ggcatcctt gtcaacacgc tgagcatggc cgtggagtac catgagcagc
2701 ccgaggagct gactaatgt ctggagatca gcaacatgt gttaccagc atgtttgcc
2761 tggagatgct gctgaagctg ctggcctgcg gccctctgg ctacatccg aaccgtaca
2821 acatcttca cggcatcatc gtgttca gctctggga gatcgtggg caggcggacg

FIG. 19B

2881 gtggcttgtc tgtgctgcgc accctccggc tgcctcgtgt gctgaagctg gtgcgcttc
 2941 tgccagccct gcggcgccag ctctgtgtgc tggtaagac catggacaac gtggctacct
 3001 tctgcacgt gctcatgctc ttcatttca tcttcagcat cctgggcatg caccttttcg
 3061 gctgcaagt cagcctgaag acagacaccg gagacaccgt gcctgacagg aagaacttcg
 3121 actccctgct gtgggccatc gtcaccgtgt tccagatcct gaccagggag gactggaaacg
 3181 tggctcgtga caacggcatg gcctccacct cctcctgggc cgcctctac ttcgtggccc
 3241 tcatgacct cggcaactat gtgctctca acctgctggt ggccatcctc gaggaggct
 3301 tccaggcgga ggccgatgcc aacagatccg acacggacga ggacaagacg tcggtccact
 3361 tcgaggagga cttccacaag ctacagagaac tccagaccac agagctgaag atgtgtccc
 3421 tggccgtgac cccaacggg cacctggagg gacgaggcag cctgtccct cccctcatca
 3481 tgtgcacagc tgccacgccc atgcctaccc ccaagagctc accattcctg gatcgacccc
 3541 ccagcctccc agactctcgg cgtggcagca gcagctccgg ggaccggcca ctgggagacc
 3601 agaagcctcc ggccagcctc cgaagtctc cctgtgccc ctggggcccc agtggcgct
 3661 ggagcagccg gcgctccagc tggagcagcc tgggcccgtc cccagcctc aagcgcccg
 3721 gccagtgtgg ggaacgtgag tccctgctgt ctggcgaggg caagggcagc accgacgacg
 3781 aagctgagga cggcagggcc gcgcccggc cccgtgccac cccactgcgg cgggcccagt
 3841 ccttgagccc acggccctg cggccggccg cctcccgcc taccaagtgc cgcgatcgcg
 3901 acgggcaggt ggtggccctg cccagcgact tcttctgctg catcgacagc caccgtgagg
 3961 atgcagccga gcttgacgac gactcggagg acagctgctg cctccgcctg cataaagtgc
 4021 tggagcccta caagccccag tgggtccgga gccgcgaggc ctgggcccct tacctttct
 4081 cccacagaa ccggttccgc gtctcctgcc agaaggtcat cacacacaag atgtttgatc
 4141 acgtgttct cgtcttcac ttcctcaact gcgtacccat cgcctggag aggcctgaca
 4201 ttgaccccg cagcaccgag cgggtcttcc tcagcgtctc caattacatc ttcacggcca
 4261 tctcgtggc ggagatgatg gtgaaggtgg tggccctggg gctgctgtcc ggcgagcacg
 4321 cctacctga gagcagctgg aacctgctgg atgggctgct ggtgctggtg tccctggtgg
 4381 acattgtct ggccatggcc tcggctggtg gcgccaagat cctgggtgtt ctgcgcgtgc
 4441 tgcgtctgt gcggaccctg cggcctctaa gggcatcag ccggggcccc ggcccaagc
 4501 tgggtgtgga gacgctgata tcgtcgtca ggccattgg gaacatcgtc ctcatctgt
 4561 gcgcttctt catcatttt ggcatctgg gtgtgcagct ctcaaaggg aagtctact
 4621 actgcgaggg cccgacacc aggaacatct ccaccaaggc acagtgcgg gccgcccact
 4681 accgtgggt gcgacgcaag tacaacttcg acaacctggg ccaggccctg atgtcgtgt
 4741 tcgtgctgt atccaaggat gtaggggtga acatcatgta cgacgggctg gatccgtgg
 4801 gtgtcgacca gcagcctgtg cagaaccaca acccctggat gctgctgtac ttcatctct
 4861 tctgtctcat cgtcagctc ttcgtgctca acatgttctg gggcgtcgtg gtcgagaat
 4921 tccacaagt ccggcagcac caggaggcgg agggaggcgg gcggcgagag gagaagcggc
 4981 tgcggcgct agagaggagg cgcaggagca cttccccag ccagaggcc cagcgcggc
 5041 cctactatgc cgactactc cccacgcgc gctccattca ctgctgtgc accagccact
 5101 atctcagct cttcatcacc ttcatctct gtgtcaact catcacatg tccatggagc
 5161 actataacca acccaagtgc ctggacgagg cctcaagta ctgcaactac gtcttacc
 5221 tcgtgttgt cttcagggt gcactgaagc tggtagcatt tgggttccgt cgttttca
 5281 aggacaggtg gaaccagctg gacctggcca tcgtgtgtc gtcactcatg ggcacacgc
 5341 tggaggagat agagatgagc gccgcgtgc ccatcaacc caccatcatc cgcacatgc
 5401 gcgtgcttc cattgcccgt gtgtgaagc tgtgaagat ggctacgggc atgcgcgcc
 5461 tgcgggacac tgtgtgcaa gctcctccc aggtggggaa cctgggctt ctttcatgc
 5521 tctgttttt tatctatgt gcgctgggag tggagctgt cgggaggctg gagtgcagt
 5581 aagacaaccc ctgcgagggc ctgagcaggc acgccacct cagcaacttc ggcatggcct
 5641 tctcacact gtccgcctc tccacgggg acaactgaa cgggacatg aaggacacgc

FIG. 19C

5761 tctacttcgt gaccttcgtg ctggtggccc agttcgtgct ggtgaacgtg gtggtggccc
5821 tgctcatgaa gcacctggag gagagcaaca aggaggcacg ggaggatgcg gagctggacg
5881 ccgagatcga gctggagatg gcgcagggcc ccgggagtcg acgccgggtg gacgcggaca
5941 ggccctccctt gccccaggag agtccgggcg ccagggatgc cccaaacctg gttgcacga
6001 aggtgtccgt gtccaggatg ctctcgtgc ccaacgacag ctacatgttc aggccgtgg
6061 tgctgcctc ggcgccccac cccgcccgc tgcaggaggt ggagatggag acctatggg
6121 ccggcacccc ctgggctcc gtgctctg tgcactctcc gccgcagag tctgtgcct
6181 ccctccagat cccactggct gtgtcgtccc cagccaggag cggcgagccc ctccacgcc
6241 tgtccctcg gggcacagcc cgctcccga gtctcagccg gctgctctg agacaggagg
6301 ctgtgcacac cgattcctt gaagggaaga ttgacagccc tagggacacc ctggatcctg
6361 cagagcctgg tgagaaaacc ccggtgaggc cggtagccca ggggggctcc ctgcagtc
6421 caccacgctc cccacggccc gecagcgtcc gcactcgtaa gcatacctc ggacagcact
6481 gcgtctccag ccggccggcg gcccaggcg gagaggaggc cgaggcctcg gacccagccg
6541 acgaggaggt cagccacatc accagctccg cctgcccctg gcagcccaca gccgagcccc
6601 atggccccga agcctctccg gtggccggcg gcgagcggga cctgcgcagg ctctacagcg
6661 tggacgtca gggcttctg gacaagccgg gccgggcaga cgagcagtg gggccctcg
6721 cggagctggg cagcggggag cctggggagg cgaaggcctg gggccctgag gccgagcccg
6781 ctctgggtg gcgcagaaag aagaagatga gccccctg catctcgtg gaacccctg
6841 cggaggacga gggctctcg cggccctccg cggcagaggc cggcagcacc acctgaggc
6901 gcaggacccc gtctgtgag gccacgcctc acagggactc cctggagccc acagagggt
6961 caggcgccgg gggggaccct gcagccaagg gggagcgtg gggccaggcc tctgcccgg
7021 ctgagcacct gaccgtcccc agcttgcct ttgagccgt ggacctcggg gtcccagtg
7081 gagaccctt ctggacggt agccacagt tgacccaga atccagagct tctcttcag
7141 gggccatagt gccctggaa ccccagaat cagagcctc catgccgtc ggtgacccc
7201 cagagaagag gcgggggctg tacctcacag tccccagt tctctggag aaaccagggt
7261 cccctcagc ccccctgcc ccagggggtg gtgcagatga cccgtgtg ctgggggtt
7321 ggtgccgccc acggcttgg ccctggggtc tgggggcccc gctggggtg agggccaggc
7381 agaaccctgc atggacctg acttgggtcc cgtcgtgagc agaaaggccc ggggaggatg
7441 acggcccagg ccctggttct ctgccagcg aagcaggagt agctgccggg ccccacgagc
7501 ctccatccgt tctggtcgg gtttccga gtttgcctac cagccaggc tgtcgggca
7561 actgggtcag cctccgtca ggagagaagc cgcgtctgtg ggacgaagac cgggcacccc
7621 ccagagaggg gaaggtacca ggtgctcc ttacaggccc cgcgttgta caggacactc
7681 gctggggcc ctgtgccctt gccggcgca ggtgcagcc accgcggccc aatgtcacct
7741 tcactcacag tctgagttct tgcgcctg tcacgcctc accaccctc cctccagcc
7801 accaccctt ccgttccgt cgggcctcc cagaagcgtc ctgtactct gggagagggtg
7861 acacctact aaggggccga cccatggag taacgcgc

FIG. 19D

MTEGARAADDEVVRVPLGAPPPGPAALVGASPESPGAPGREAGERGS
ELGVSPSESPAERGAELGADEEQRVYPALAAATVFFCLGQTTRPRSWCLRLVCNPWF
EHVSMMLVIMLNCVTLMFRPCEDVECGSERCNILEAFDAFIFAFFAVEMVIKMVALGL
FGQKCYLGDTWNRLDFFIVVAGMMEYSLDGHNVSLSAIRTVRVLRLRAINRVPSMRI
LVTLLDTPMLGNVLLCFFVFFIFGIVGVQLWAGLLRNRCFLDSAFVRNNNLTLFLR
PYYQTEEGEENPFICSSRRDNGMQKCSHIPGRRELMPCTLGWEAYTQPQAEVGGAAR
NACINWNQYYNVCRSGDSNPHNGAINFDNIGYAWIAIFQVITLEGWVDIMYYVMDAHS
FYNFIYFILLIIVGSFFMNLCLVVIATQFSETKQRESQLMREQRARHLSNDSTLASF
SEPGSCYEELLKYVGHIFRKVKRRSLRLYARWQSRWRKKVDPSAVQQGPGHRQRRAG
RHTASVHHLVYHHHHHHHHHHYHFSHSGSPRRPGPEPGACDTRLVRAGAPSPSPSGRGP
PDAESVHSIYHADCHIEGPQERARVAHAAATAAASRLRLATGLGTMNYPTILPSGVGSG
KGSTSPGPKGWAGGPPGTGGHGPLSLNSPDYKIPHVVGEGHGLQAPGHLGLSVLP
CPLSPAGTTLTCELKSCPYCTRALEDPEGELSGSESGLSDGRGVYEFTQDVRHGDRW
DPTRPPRATDTPGPGPGSPQRRRAQRAAPGEPGWMGRLWVTFSGKLRRIVDSKYFSRG
IMMAILVNTLSMGVEYHEQPEELTNALEISNIVFTSMFALEMLLKLLACGPLGYIRNP
YNIFDGIIVVISVWEIVGQADGGLSVLRTFRLRLVLKLVRLPALRRQLVVLVKTMDN
VATFCTLLMLFIFISILGMHLFGCKFSLKTDGTGDTVDRKNFDSLLWAIVTVFQILT
QEDWNVVLNGMASTSSWAALYFVALMTFGNYVLFNLLVAILVEGFQAEGDANRSDTD
EDKTSVHFEEDFHKLRELQTTELKMCSLAVTPNGHLEGRGSLSPPLIMCTAATPMPTP
KSSPFLDAAPSLPDSRRGSSSSGDPPLGDQKPPASLRSSPCAPWGPSGAWSSRRSSWS
SLGRAPSLKRRGQCGERESLLSGEGKGSTDDAEDGRAAPGPRATPLRRAESLDPRPL
RPAALPPTKCRDRDGQVVALPSDFFLRIDSHREDAEELDDSDSCCLRLHKVLEPYK
PQWCRSREAWALYLFSPQNRFRVSCQKVITHKMFHDHVVLVFIFLNCVTIALERPDIDP
GSTERVFLSVSNYIFTAIFVAEMMVKVVALGLLSGEHAYLQSSWNLLDGLLVLSLVD
IVVAMASAGGAKILGVLRVLRLRLTLRPLRVISRAPGLKLVVETLISSLRPIGNVLI
CCAFFIIFGILGVQLFKGKFYYCEGPDTRNISTKAQCRAAHYRWVRRKYNFDNLGQAL
MSLFVLSSKDGWVNIMYDGLDAVGVDQQPVQNHNPWMLLYFISFLLIVSFFVLNMFVG
VVVENFHKCRQHQAEEARRREEKRLRLERRRRSTFPSPEAQRRPYADYSPTRRSI
HSLCTSHYLDLFTITFCVNVITMSMEHYNQPKSLDEALKYCNVFTTVFVFEAALKL
VAFGFRFFKDRWNQLDLAIVLLSLMGITLEEIEMSAALPINPTIIRIMRVLRIARVL
KLLKMATGMRALLDTVVQALPQVGNLGLLFMLLFFIYAALGVELFGRLECEDNPCEG
LSRHATFSNFGMAFLTLFRVSTGDNWNGIMKDTLRECSREDKHCLSYLPALSPVYFVT
FVLVAQFVLVNVVVAVLMKHLEESNKEAREDAELDAEIELEMAQGPGSARRVDADRP
LPQESPGARDAPNLVARKVSVRMLSLPNDSYMFRPVVPASAPHRPLQEVEMETYGA
GTPLGSVASVHSPPAESCASLQIPLAVSSPARSGEPLHALSPRGATARSPSLRLLCRQ
EAVHTDSLEGKIDSPRDTLDAEPGEKTPVRPVTQGGSLQSPRSPRPASVRTRKHTF
GQHCVSSRPAAPGGEEAEASDPADEEVSHITSSACPWQPTAEPHGPEASPVAGGERDL
RRLYSVDAQGFLDKPGRADQWRPSAELSGEPGEAKAWGPEAEALGARRKKKMSPP
CISVEPPAEDEGSARPSAAEGGSTTLRRRTPSCEATPHRDSLEPTEGSGAGGDPAAKG
ERWGQASCRAEHLTVPSFAFEPLDLGVPSGDPFLDGSHSVTPESRASSSGAIVPLEPP
ESEPPMPVGDPPPEKRRGLYLTVPQCPLKPGSPSATPAPGGGADDPV

FIGURE 20A

1 gcggcggcgg ctgcggcggg gggggccgggc gaggtccgct gcgggtcccgg cggctccgtg
61 gctgctccgc tctgagcgcc tggcgccccc cgcgccctcc ctgccggggc cgtggggccg
121 gggatgcacg cggggcccg gagccatggt ccgcttcggg gacgagctgg gcggccgcta
181 tggaggcccc ggcggcgagg agcggggccc gggcgggggg gccggcgggg cggggggccc
241 ggggtcccgg gggctgcagc ccggccagcg ggtcctctac aagcaatcga tcgcgcagcg
301 cgcgcggacc atggcgctgt acaaccccat ccggtcaag cagaactgct tcaccgtcaa
361 ccgctcgtc ttcgtctca gcgaggacaa cgtcgtccg aaatacgcga agcgcacac
421 cgagtggcct ccattcgagt atatgatcct ggccaccatc atcgccaact gcacgtgct
481 ggccctggag cagcacctcc ctgatggga caaacgccc atgtccgagc ggctggacga
541 cacggagccc tatttcacg ggaatctttg ctgcaggca gggatcaaaa tcacgtct
601 gggctttgct ttccacaagg gctcttacct gcggaacggc tggaaactga tggacttct
661 ggtcgtcctc acagggaacc ttgccacggc tggaaactga ttccactgc gaacactgag
721 ggtgtgctgt gtgtgaggc ccctgaagct ggtgtctggg attccaagtt tgcaggtggt
781 gctcaagtc atcatgaagg ccatgggtcc actcctgcag attgggtgc ttctctct
841 tgccatcctc atgtttgcca tcattggcct ggagttctac atgggcaagt tccacaaggc
901 ctgtttcccc aacagcaçag atgcggagcc cgtgggtgac ttccctgtg gcaaggaggc
961 cccagcccgg ctgtgcgagg gcgacactga gtgcgggag tactggccag gaccaact
1021 tggcatcacc aactttgaca atatcctgtt tgccatctg acggtgtcc agtgcacac
1081 catggaggggc tggactgaca tctctataa tacaacgat gcggccggca acacctggaa
1141 ctggctctac ttcacccctc tcacatcat cggtccttc ttcacgtca acctggtgct
1201 gggcgtgctc tcgggggagt ttccaagga gcgagagagg gtggagaacc gccgcgcctt
1261 cctgaagctg cggcgccagc agcagatcga gcgagagctc aacgggtacc tggagtggat
1321 cttcaaggcg gaggaagtca tgctggccga ggaggacagg aatgcagagg agaagtcccc
1381 ttggacgtg ctgaagagag cggccaccaa gaagagcaga aatgacctga tccacgcaga
1441 ggaggagag gaccggttg cagatctctg tgctgttga tccccctc cccgcgccag
1501 cctcaagagc gggaagacag agagctcgtc atacttccg aggaaggaga agatgttccg
1561 gtttttacc cggcgcatgg tgaaggctca gagcttctac tgggtggtgc tgtgcgtggt
1621 ggccctgaac acactgtgtg tggccatggt gcattacaac cagccgcggc ggcttaccac
1681 gaccctgtat ttgcagagt ttgtttcct gggctcttc ctacagaga tgccttgaa
1741 gatgatggc ctggggccca gaagctactt ccggtcctcc ttcaactgct tcgactttg
1801 ggtcatcgtg gggagcgtct tgaagtgtt ctggcgccg atcaagccgg gaagctcctt
1861 tgggatcagt gtgtgcggg ccctccgct gctgaggatc ttcaaagtca cgaagtactg
1921 gagctccctg cggaacctgg tgggtccct gctgaactcc atgaagtcca tcatcagcct
1981 gctctcttg ctctcctgt tcattgtggt ctgcgccctg ctggggatgc agctgtttg
2041 gggacagttc aacttccagg atgagactcc cacaaccaac ttcgacacct tcctgccgc
2101 catcctact gtcttcaga tctgacggg agaggactgg aatgcagtga tgtatcacgg
2161 gatcgaatc caaggcgcg tcagcaaagg catgtctc tcttttact tcattgtcct
2221 gacactgtc gaaactaca ctctgtgaa tgtcttctg gccatcgtg tggacaacct
2281 ggccaacgcc caagagctga ccaaggatga agaggagatg gaagaagcag ccaatcagaa
2341 gcttctctg caaaaggcca aagaagtggc tgaagtcag cccatgtctg ccggaacat
2401 ctccatgcc gccaggcagc agaactcggc caaggcgcg tcggtgtggg agcagcgggc
2461 cagccagcta cggctgcaga acctgcgggc cagctgcgag gcgtgtaca gcgagatgga
2521 ccccgaggag cggctgcgt tcgccactac gcgccactg cggcccgaca tgaagacga
2581 cctggaccgg ccgctggtg tggagctggg ccgacgagc gcgcggggg ccgtgggagg
2641 caaagccga cctgaggctg cggaggcccc caggggcgtc gacctccgc gcaggacca
2701 ccggcacgc gacaaggaca agaccccgcc ggcgggggac caggaccgag cagaggcccc
2761 gaaggcggag agcggggagc ccggtgccc gaggagcgg ccgcggccgc accgcagcca
2821 cagcaaggag gccgcggggc ccccgaggc gcggagcag cgcggccgag gccagggccc

FIGURE 20B

2881 cgagggcggc cggcggcacc accggcgcgg ctccccggag gaggcggccg agcggggagcc
 2941 ccgacgccac cgcgcgcacc ggcaccagga tccgagcaag gagtgcgccc gcgccaaggg
 3001 cgagcggcgc gcgcggcacc gcggcggccc ccgagcgggg ccccgggagg cggagagcgg
 3061 ggaggagccg gcgcggcggc accggggccc gcacaaggcg cagcctgctc acgaggctgt
 3121 ggagaaggag accacggaga aggaggccac ggagaaggag gctgagatag tggaaagcga
 3181 caaggaaaag gagctccgga accaccagcc ccgggagcca cactgtgacc tggagaccag
 3241 tgggactgtg actgtgggtc ccatgcacac actgcccagc acctgtctcc agaaggtgga
 3301 ggaacagcca gaggatgcag acaatcagcg gaacgtcact cgcattggga gtcagcccc
 3361 agaccgaac actattgtac atatccagt gatgtgacg ggccctcttg gggaagccac
 3421 ggtcgttccc agtgtaacg tggacctgga aagccaagca gaggggaaga aggaggtgga
 3481 agcggatgac gtgatgagga gcggccccc gcctatctc ccatacagct ccattgtctg
 3541 tttaagcccc accaactgc tccgccgtt ctgccactac atcgtgacca tgaggtaact
 3601 cgagggtgct attctcgtgg tcctgcctt gagcagcatc gccctggctg ctgaggaccc
 3661 agtgcgcaca gactcgccca ggaacaacgc tctgaaatac ctggattaca ttttactgg
 3721 tgtctttacc ttgagatgg tgataaagat gatcgacttg ggactgctgc ttcaccctgg
 3781 agcctatttc cgggacttgt ggaacattct ggacttcaat gtggtcagtg gcgccctgg
 3841 ggcgtttgct ttctcaggat ccaagggaa agacatcaat accatcaagt ctctgagagt
 3901 ccttcgtgic ctgcggcccc tcaagaccat caaacggctg ccaagctca aggtgtgtt
 3961 tgactgtgtg gtgaactccc tgaagaatgt cctcaacatc ttgattgtct acatgtctt
 4021 catgttcata ttgccgtca ttgcggtgca gctcttcaa gggaagtitt tctactgcac
 4081 agatgaatcc aaggagctgg agagggactg caggggctag tatttggtat atgagaagga
 4141 ggaagtggaa gctcagccca ggcagtggaa gaaatacagc ttctactagc acaatgtgct
 4201 ctgggctctg ctgacgtgtg tcacagtgc cacgggagaa ggctggccca tgggtctgaa
 4261 aactccgtg gatgccacct atgaggagca ggtccaagc cctgggtacc gcatggagct
 4321 gtccatcttc tacgtggtct acittgtgtt ttcccttc ttctctgca acatcttgt
 4381 ggctttgatc atcatcacct tccaggagca gggggacaag gtgatgtctg aatgcagcct
 4441 ggagaagaac gagagggtt gcattgactt cgccatcagc gccaaacccc tgacacggt
 4501 catgccccaa aaccggcagt cgttccagta taagacgtgg acatttgttg tctccccgcc
 4561 ctttgaatac ttcatcatgg ccattatagc cctcaacact gtgtgtctga tgatgaagt
 4621 ctatgatgca ccctatgagt acgagctgat gctgaaatgc ctgaacatcg tgttcacatc
 4681 catgttctcc atggaatcgc tctgaagat catcgcttt gggtgtctga actatttcag
 4741 agatgcctgg aatgtcttg actttgtcac tgtgttgga agtattactg atatttagt
 4801 aacagagatt gcggaaacga acaatttcat caacctcagc ttcctccgc tcttcgagc
 4861 tgcgcggctg atcaagctgc tccgccaggg ctacaccatc cgcacctgc tgtggacct
 4921 tgtccagtcc ttcaaggccc tgcctacgt gtgtctgctc attgcatgc tgttctcat
 4981 ctaccgatc atcggcagc aggtgtttgg gaatattgcc ctggatgatg acaccagcat
 5041 caaccggcac acaacttcc ggacgtttt gcaagccctg atgtgtgtg tcaggagcgc
 5101 cacgggggag gcctggcacg agatcatgct gtctgcctg agcaaccagg cctgtgatga
 5161 gcaggccaat gccaccgagt gtggaagtga ctttgcctac ttctacttcg tctcttcat
 5221 ctctctgic tctttctga tgtgaacct cttgtggct gtgatcatg acaatttga
 5281 gtacctcag cgggactctt ccatcctagg tctcaccac ttggatgagt tcatccgggt
 5341 ctgggctgaa tacgaccgg ctgcgtgtgg gcgcatcagt tacaatgaca tgtttgagat
 5401 gctgaacac atgtccccg ctctggggct gggaagaaa tgcctgctc gattgtctta
 5461 caagcgctg gttcgatga acatgccat ctccaacgag gacatgactg ttcacttcac
 5521 gtccacgctg atggccctca tccggacggc actggagatc aagctggccc cagctgggac
 5581 aaagcagcat cagtgtgacg cggagttag gaaggagatt tccgttgtgt gggccaatct

FIG. 20C

5641 gccccagaag actttggact tgctggtacc accccataag cctgatgaga tgacagtggg
5701 gaaggtttat gcagctctga tgatattga cttctacaag cagaacaaaa ccaccagaga
5761 ccagatgcag caggctcctg gaggcctctc ccagatgggt cctgtgtccc tgtccaccc
5821 tctgaaggcc accctggagc agacacagcc ggctgtgctc cgaggagccc gggttttcct
5881 tcgacagaag agttccacct cctcagcaa tggcggggcc atacaaaacc aagagagtgg
5941 catcaaagag tctgtctcct ggggcactca aaggaccag gatgcacccc atgaggccag
6001 gccacccctg gagcgtggcc actccacaga gatccctgtg gggcgggtcag gagcactggc
6061 tgtggacgtt cagatgcaga gcataaccg gaggggccct gatggggagc cccagcctgg
6121 gctggagagc cagggtcgag cggcctccat gcccgcctt gcggccgaga ctacagccgt
6181 cacagatgcc agcccatga agcgtccat ctccacgtg gccagcggc cccgtgggac
6241 tcattttgc agcaccacc cggaccgcc accccctagc caggcgtctg cgcaccacca
6301 ccaccaccgc tgccaccgcc gcaggagacag gaagcagagg tccctggaga agggggccag
6361 cctgtctgcc gatatggatg gcgcaccaag cagtgtgtg gggccggggc tgccccggg
6421 agagggggcct acaggctgcc ggcgggaacg agagcgccgg caggagcggg gccgggtcca
6481 ggagcggagg cagccctcat cctcctcctc ggagaagcag cgcttctact cctgcgaccg
6541 ctttgggggc cgtgagccc cgaagcccaa gccctcctc agcagccacc caacgtcgcc
6601 aacagctggc caggagccgg gacccaccc acagggcagt ggttccgtga atgggagccc
6661 cttgctgtca acatctggtg ctacacccc cggcgcgggt gggcgggagc agtccccca
6721 gacgcccctg actccccgcc ccagcatcac ctacaagacg gccaactcct caccatcca
6781 ctccgccggg gctcagacca gccctcctgc cttctccca ggcgggtca gccgtgggt
6841 ttccgaacac aacgccctgc tgcagagaga cccctcagc cagccccgg cccctggctc
6901 tcgaattggc tctgacctt acctggggca gcgtctggac agtgaggcct ctgtccacgc
6961 cctgcctgag gacacgtca cttcgagga ggctgtggc accaactcgg gccgtcctc
7021 caggacttcc tacgtgtcct cctgacctc ccagtctac cctctccgcc gcgtgccccaa
7081 cggttaccac tgcacctgg gactcagtc gggtgggcga gcacggcaca gctaccacca
7141 ccctgaccaa gaccactgg gctagtgca ccgtgaccg tcagacgcct gcatgcaga
7201 ggcgtgtgtt ccagtggatg agttttatca tccacacggg gcagtcggcc ctgggggag
7261 gccctgcca ccttggtgag gctcctgtgg cccctccctc cccctcctc cctctttac
7321 tctagacgac gaataaagcc ctgttgctt agtgtacgta ccgc

FIG. 20D

MVRFGDELGGRYGGPGGGERARGGGAGGAGGPGPGGLQPGQRVL
 YKQSIQRARTMALYNPIPVKQNCFTVNRSFLVFSEDNVVRKYAKRITWPPFEYML
 ATIANCIVLALEQHLPGDKTPMSERLDDTEPYFIGIFCFEAGIKIILGFVFHKGS
 YLRNGWNVMDFVVVLTGILATAGTDFDLRTLRAVRVLRPLKLVSGIPSLQVVLKSMK
 AMVPLLQIGLLLFFAILMFAIIGLEFYMGKFHKACFPNSTDAEPVGDFFCGKEAPARL
 CEGDTECREYWP GPNFGITNFDNLFAILTVFQCITMEGWTDILYNTNDAAGNTWNWL
 YFIPLIIGSFFMLNLVLGVLSGEFAKERERVENRRAFLKLRQQQIERELNGYLEWI
 FKAEEVMLAEEDRNAEEKSPLDVLKRAATKKSRNDLIHAEEGEDRFADLCAVGSPFAR
 ASLKSGKTESSYFRRKEKMFRFFIRRMVKAQSFYVWVLCVVALNTLCVAMVHYNQPR
 RLTTTLYFAEFVFLGLFLTEMSLKMYGLGPRS YFRSSFNCDFGVIVGSVFEVWAAI
 KPGSSFGISVLRALRLRIFKVTKYWSSLRNLVVSLLNSMKSIISLLFLFLFVVFVFA
 LLGMQLFGGQFNFDPTTNTFDTPFAAILTVFQILTGEDWNAV MYHGIESQGGVSKG
 MFSSFYFVLTFLGNYTLLNVFLAIAVDNLANAQELTKDEEEMEEAANQKLALQKAKE
 VAEVSPMSAANISIAARQQNSAKARSVWEQRASQLRLQNLRASCEALYSEMDPEERLR
 FATTRHLRPDMKTHLDRPLVVELGRDGARGPVGGKARPEAAEAPEGVDP PRRHHRHRD
 KDKTPAAGDQDRAEAPKAESGEPGAREERPRPHRSKSKEAAGPPEARSERGRGPGE
 GRRHHRGSPPEAAEREPRRHRAHRHQDPSKECAGAKGERRARHRGGPRAGPREAESG
 EEPARRHRARHKAQPAHEAVEKETTEKEATEKEAEIVEADKEKELRNHQPREPHCDLE
 TSGTVTVGPMHTLPSTCLQKVEEQPEDADNQNRNVTRMGSQPPDPNTIVHIPVMLTGPL
 GEATVVP SGNV DLESQAEGKKEVEADDVMRSGPRPIVPYSSMFCLSPTNLLRRFCHYI
 VTMR YFEV VILV VIALSSIALAAEDPVRTDSPRNNALKYLDYIFTGVFTFEMVIK MID
 LGLLLHPGAYFRDLWNILDFIVVSGALVAFAFSGSGKGDINTIKSLRVLRVLRPLKTI
 KRLPKLKA VFDCV VNSLKNVNLIVYMLFMFIFAVIAVQLFKGKFFYCTDESKELER
 DCRGQYLDYEKEEVEAQPRQWKYDFHYDNVLWALLTLFTVSTGEGWPMVLKHSVDAT
 YEEQGSPGYRMELSIFYVVYFVFPFFFNIFVALIITFQE QGDKVMSECSLEKNE
 RACIDFAISAKPLTRYMPQNRQSFQYKTWTFVVSPPFEYFIMAMIALNTVVLMMKFYD
 APYEYELMLKCLNIVFTSMFSMECVLKIIAFGVNLNYFRDAWNVDFVTVLGSITDILV
 TEIAETNNFINLSFLRLFRAARLIKLLRQGYTIRILLWTFVQSFKALPYVCLLIAMLF
 FTYAIIGMQVFGNIALDDDT SINRHNNFRFTLQALMLLFRSATGEAWHEIMLSCLSNQ
 ACDEQANATECGSDFAYFYFVSFIFLCSFLMLNLFVA VMNDNFEYLTRDSSILGPHHL
 DEFIRVWAEYDPAACGRISYNDMFEMLKHMSPPLGLGKKCPARVAYKRLVRMNMPI SN
 EDMTVHFTSTLMALIRTALEIKLAPAGTKQHQCDAELRKEISVWVANLPQKTLDLLVP
 PHKPDEMTVGKVYAALMIFDFYKQNKTT RDQMQQAPGGLSQMGVSLFHPLKATLEQT
 QPAVLRGARVFLRQKSSTLSNGGAIQNQESGIKESVSWGTQRTQDAPHEARPPLERG
 HSTEIPVGRSGALAVDVQMQSITRRGPDGEPQPGLESQGRAASMPRLAAETQPVT DAS
 PMKRSISTLAQRPRGTHLCSTTPDRPPPSQASSHHHHHRCHRRRDRKQRSLEKGPSLS
 ADMDGAPSSAVGPGLPPGEGPTGCRERERRQERGRSQERRQPSSSSSEKQRFYSCDR
 FGGREPPKPKPSLSSHPTSPTAGQEPGHPQGSGSVNGSPLLSTSGASTPGRGRRQL
 PQTPLTPRPSITYKTANSSPIHFAGAQTSLPAFSPGRLSRGLSEHNALLQRDPLSQPL
 APGSRIGSDPYLGQRLDSEASVHALPEDTLTFEEAVATNSGRSSRTSYVSSLTSQSH
 LRRVPNGYHCTLGLSSGGRARHSYHHPDQDHC

FIGURE 21A

1 gcggcggcgg ctgcggcggg gggggcgggc gaggtccgct gcggtcggcg cggctccgtg
61 gctgtccgc tctgagcgcc tggcgcggcc cgcgccctcc ctgcggggcg cgttgggcgg
121 gggatgcacg cggggcccg gagccatggt ccgcttcggg gacgagctgg gcggccgcta
181 tggagggccc ggcggcggag agcggggccc gggcggcggg gccggcgggg cggggggccc
241 ggggtccggg gggctgcagc ccggccagcg ggtctctac aagcaatga tcgcgcagcg
301 cgcgcggacc atggcgctgt acaaccccat ccggtcaag cagaactgct tcaccgtcaa
361 ccgctcgctc ttctttca gcgaggacaa cgtctccgc aaatacgcga agcgcatcac
421 cgagtggcct ccattcgagt atatgatcct ggccaccatc atcgcaact gcctgtgct
481 ggccctggag cagcacctcc ctgatgggga caaacgccc atgtccgagc ggctggacga
541 cacggagccc tatttcacg ggatctttg ctgcaggca gggatcaaaa tcctgctct
601 gggctttgtc ttccacaagg gctcttacct gcggaacggc tggaaactga tggacttct
661 ggtctgctc acagggatcc ttgccacggc tggaaactgac ttgcacctgc gaacactgag
721 ggtctgctc gtgctgaggc cctgaagct ggtgtctggg attccaagtt tgcaggtggt
781 gctcaagtc atcatgaagg ccatggtcc actcctgcag attgggctgc ttcttctt
841 tgccatctc atgtttgca tcattggcct ggagtctac atgggcaagt tccacaaggc
901 ctgttcccc aacagcacag atgcggagcc cgtgggtgac ttccctgtg gcaaggaggc
961 cccagcccg ctgtgcagg gcgacactga gtccggggg tactggccag gacccaact
1021 tggcatcac aactttgaca atactctgt tggcatctg acggtgtcc agtgcacac
1081 catggagggc tggactgaca tcctctataa tacaacgat gcggccggca acactggaa
1141 ctggtctac ttatccctc tcattcatc atggtcttc ttcatgtca acctggtgt
1201 gggcgtgtc tcgggggagt ttgcaagga gcgagagagg gtggagaacc gccgcgctt
1261 cctgaagctg cgcggcagc agcagatga gcgagagctc aacgggtacc tggagtggat
1321 ctcaaggcg gaggaagtca tgtggccga ggaggacagg aatgcagagg agaagtcgcc
1381 ttggacgtg ctgaagagag cggccacca gaagagcaga aatgacctga tccacgaga
1441 ggaggagag gaccggttg cagatctctg tctgttga tccccttcg ccgcgccag
1501 cctcaagagc gggaagacag agagctcgtc tttctccg aggaaggaga agatgtccg
1561 gtttttatc cgcgcgatg tgaaggctca gagcttctc tgggtgtgc tgtgctggt
1621 ggccctgaac acactgtgtg tggccatggt gcattacaac cagcccggc ggcttaccac
1681 gacctgtat ttgcagagt ttgtttctt ggtctcttc ctacagaga tgcctgaa
1741 gatgatggc ctggggcca gaagctactt ccggtctcc ttcaactgct tcgactttg
1801 ggtcatcgtg gggagcgtct tgaagtgtt ctgggcggcc atcaagccgg gaagctctt
1861 tgggatcagt gtgtcggg cctccgct gctgaggatc ttcaagtca cgaagtactg
1921 gagctccctg cggaaactgg tgggtctct gctgaactcc atgaagtcca tcacagcct
1981 gctcttctg ctctctgt tcattgtgt ctccgctg ctggggatgc agctgtttg
2041 gggacagtc aacttcagg atgagactcc cacaaccaac ttgacacct tcctgccgc
2101 cactctact gtctccaga tctgacgg agaggactgg aatgcagtga tttatcacg
2161 gatgaatcg caaggcggc tcagcaaagg catgttctg tcctttact tcattgtct
2221 gacactgtc ggaactaca ctctgtgaa tgtttctg gccatcgtg tggacaacct
2281 ggccaacgcc caagagctga ccaaggatga agaggagatg gaagaagcag ccaatcagaa
2341 gctgtctg caaaaggcca aagaagtggc tgaagtcagc ccatgtctg ccgcgaacat
2401 ctccatgcc gccagcagc agaactcggc caaggcgcgc tgggtgtgg agcagcgggc
2461 cagccagcta cggctgcaga acctgcgggc cagctgcgag gcgctgtaca gcgagatgga
2521 cccgaggag cggctgcgt tcgccactac gcgccacctg cggcccgaca tgaagacga
2581 cctggaccgg ccgtggtgg tggagctggg ccgcgacggc gcgcggggg ccgtgggagg
2641 caaagcccga cctgaggctg cggaggcccc cgaggcgct gacctccgc gcaggacca
2701 ccggcaccgc gacaaggaca agaccccg gcggggggac caggaccgag cagaggcccc
2761 gaaggcggag agcggggagc ccggtgccc ggaggagcgg ccgcggccgc accgagcca

FIGURE 21B

2821 cagcaaggag gccgcggggc ccccgagggc gcggagcgag cgcggccgag gccagggccc
2881 cgagggcggc cggcgggcacc accggcgcg cccccggag gaggcggccg agcgggagcc
2941 ccgacgccac cgcgcgcacc ggaccaggga tccgagcaag gaggcgccg gcgccaaggg
3001 cgagcggcg gcgcggcacc gcggcgggcc ccgagcgggg ccccgggagg cggagagcgg
3061 ggaggagccg gcgcggcgcc accgggccc gcacaaggcg cagcctgctc acgaggctgt
3121 ggagaaggag accacggaga aggaggccac ggagaaggag gctgagatag tggaaagccga
3181 caaggaaaag gagctccgga accaccagcc ccgggagcca cactgtgacc tggagaccag
3241 tgggactgtg actgtgggtc ccatgcacac actgcccagc acctgtctcc agaagggtga
3301 ggaacagcca gaggatgcag acaatcagcg gaacgtcact cgcattgggca gtcagcccc
3361 agaccgaac actattgtac atatccagt gatgtgacg ggccctcttg gggaagccac
3421 ggtcgttccc agtggttaac tggacctgga aagccaagca gaggggaaga aggaggtgga
3481 agcggatgac gtgatgagga gcggccccc gcctatcgtc ccatcagct ccatgtctg
3541 tttaagcccc accaacctgc tccgccgtt ctgcactac atcgtgacca tgaggtactt
3601 cgagggtgtc attctctgtg tcatcgccct gagcagcacc gccctggctg ctgaggacc
3661 agtcgcaca gactcgcca ggaacaacgc tctgaaatac ctggattaca ttttactgg
3721 tgtcttacc ttgagatgg tgataaagat gatcgactg ggactgctgc ttcacctgg
3781 agcctattc cgggacttgt ggaacattct ggacttcatt tgggtcagt gcgccctggt
3841 ggcgtttgct tctcaggat ccaaaggga agacatcaat accatcaagt cctgagagt
3901 cctcgtgtc ctgcggccc tcaagaccat caacggctg ccaagctca aggtgtgtt
3961 tgactgtgtg tgaactccc tgaagaatgt cctcaacatc ttgattgtct acatgctctt
4021 catgtcata ttgccgca tgcgggtgca gctctcaaa gggaagttt tctactgcac
4081 agatgaatcc aaggagctgg agagggactg caggggctag tatttgatt atgagaagga
4141 ggaagtggaa gtcagccca ggcagtggaa gaaatacgac ttctactacg acaatgtgt
4201 ctgggctctg ctgacgtgt tcacagtgc cacgggagaa ggctggccca tgggtctgaa
4261 aactccgtg gatgccacct atgaggagca ggtccaagc cctgggtacc gcattggagt
4321 gtccatctc tacgtgtgt actttgtgt ctttccctc ttctctgca acatcttgt
4381 ggctttgat atcatcacct tccaggagca gggggacaag gtgatgtctg aatgcagcct
4441 ggagaagaac gagagggtt gcattgactt cgccatcagc gccaaacccc tgacacgga
4501 catgccccaa aaccggcagt cgttcagta taagacgtg acatttgtg tctcccgcc
4561 cttgaatac tcatcatgg ccatgatag cctcaacact tgggtgctga tgatgaagt
4621 ctatgatga ccctatgag acgagctgat gctgaaatgc ctgaacatc tttcacatc
4681 catgttctc atggaatgc tgcgaagat catgcctt gggtgctga actattcag
4741 agatgcctg aatgtcttg actttgtac tgtgttgga agtattactg atatttagt
4801 aacagagatt gcggaacga acaattcat caacctcagc ttctccgcc tcttcgagc
4861 tgcgcggctg atcaagctg tccgccagg ctacaccatc cgcctcctg tgtggacct
4921 tgtccagtc tcaaggccc tgcctacgt gtgtctgctc attgccatg tgttctcat
4981 ctacgccatc atggcatgc aggtgtttg gaattattgc ctggatgatg acaccagcat
5041 caaccgccac aacaactcc ggacgtttt gcaagccctg atgtgctgt tcaggagcgc
5101 cacgggggag gcctggcacg agatcatgct gtctgcctg agcaaccagg cctgtgatga
5161 gcaggccaat gccaccgagt gtggaagtga cttgcctac ttctactcg tctcctcat
5221 cttctgtgc tctttctga tgtgaacct cttgtggt gtgatcatg acaatttga
5281 gtacctcag cgggactct ccatcctagg tctcaccac ttggatgagt tcatccgggt
5341 ctgggctgaa tacgaccgg ctgcgtgtg gcgcatcagt tacaatgaca tgttgagat
5401 gctgaaacac atgccccgc cctggggct gggaagaaa tgcctgctc gattgtcta
5461 caagcgctg gttcgcatg acatgccat ctccaacgag gacatgactg ttacttcac
5521 gtccacgctg atggccctca tccggacggc actggagatc aagctggccc cagctgggac

FIG. 21C

5581 aaagcagcat cagtgtgacg cggagttgag gaaggagatt tccgttgtgt gggccaatct
5641 gccccagaag actttggact tgctggtagc acccataag cctgatgaga tgacagtggg
5701 gaagggttat gcagctctga tgatattga ctctacaag cagaacaaaa ccaccagaga
5761 ccagatgcag caggctcctg gaggcctctc ccagatgggt cctgtgtccc tgtccaccc
5821 tctgaaggcc accctggagc agacacagcc ggctgtgctc cgaggagccc gggtttctct
5881 tcgacagaag agtccacct cctcagcaa tggcggggcc atacaaaacc aagagagtgg
5941 catcaaagag tctgtctct ggggcactca aaggaccag gatgcacccc atgaggccag
6001 gccacccctg gagcgtggcc actccacaga gatccctgtg gggcggtagc gagcactggc
6061 tgtggacgtt cagatgcaga gcataaccg gagggggcct gatggggagc cccagcctgg
6121 gctggagagc cagggtcgag cggcctccat gccccgctt gcggccgaga ctacagccgt
6181 cacagatgcc agcccatga agcgtccat ctccacgtg gccagcggc cccgtggagc
6241 tcatctttgc agcaccacc cggaccgcc accccctagc caggcgtcgt cgcaccacca
6301 ccaccaccgc tggcaccgcc gcaggagacag gaagcagagg tccctggaga agggggccag
6361 cctgtctgcc gatattgatg gcgcaccaag cagtgtgtg gggccggggc tgccccggg
6421 agagggggcct acaggctgcc ggcgggaacg agagcgccgg caggagcggg gccggtccca
6481 ggagcggagg cagccctcat cctctctc gcgagaagcag cgcttctact cctgcgaccg
6541 ctgtggggc cgtgagcccc cgaagccaa gccctccctc agcagccacc caacgtcgc
6601 aacagctggc caggagccgg gacccaccc acaggccggc tcagccgtgg gctttccgaa
6661 cacaacgccc tgcgcagag agacccctc agccagccc tggccctgg ctctgaatt
6721 ggctctgacc ctacctggg gcagcgtctg gacagttagg cctctgtcca cgcctgct
6781 gaggacacgc tcaatttga ggaggtgtg gccaccaact cgggcccctc ctccaggact
6841 tctacgtgt ctccctgac ctccagtct caccctctcc gccgcgtgcc caacggtac
6901 cactgcaccc tgggactcag ctgggtggc cgagcacggc acagctacca ccaccctgac
6961 caagaccact ggtgctagct gcaccgtgac cgctcagac cctgcatgca gcaggcgtgt
7021 gttccagtgg atgagttta tcatccacac ggggcagtcg gccctcgggg gaggccttgc
7081 ccaccttggg gaggtcctg tggccctcc ctccctctcc tccccttct tactctagac
7141 gacgaataaa gccctgttc ttgagtgtac gtaccgc

FIG. 21D

MVRFGDELGGRYGGPGGGERARGGGAGGAGGPGGGLQPGQRVL
YKQSIAQRARTMALYNPIPVKQNCFTVNRSLFVFSEDNVVRKYAKRITEWPPFEYML
ATHANCIVLALEQHLPDGDKTPMSERLDDTEPYFIGIFCFEAGIKIALGFVFHKG
YLRNGWNVMDFVVVLTGILATAGTDFDLRTLRAVRVLRPLKLVSGIPSLQVVLKSMK
AMVPLLQIGLLFFAILMFAIIGLEFYMGKFHKACFPNSTDAEPVGDFFPCGKEAPARL
CEGDTECREYWPGPNFGITNFDNILFAILTVFQCITMEGWTDILYNTNDAAGNTWNWL
YFIPLIIGSFFMLNLVLGVLSGEFAKERERVENRRAFLKLRRQQQIERELNGYLEWI
FKAEEVMLAEEDRNAEEKSPLDVLKRAATKKSRLNDLIHAEEGEDRFADLCAVGSPPAR
ASLKSGKTESSSYFRRKEKMFRFFIRRMVKAQSFYWVLCVVALNTLCVAMVHYNQPR
RLTTTLYFAEFVFLGLFLTEMSLKMYGLGPRS YFRSSFNCFDFGVTVGSVFVWAAI
KPGSSFGISVLRALRLRIFKVTKYWSSLRLNVVSLLSNMKSIISLLFLLFLFVFA
LLGMQLFGGQFNQDETPTTNFDTFPAAILTVFQILTGEDWNAV MYHGIESQGGVSKG
MFSSFYFTVLTFLGNYTLLNVFLAIAVDNLANAQELTKDEEEMEEAANQKLALQKAKE
VAEVSPMSAANISIAARQQNSAKARSVWEQRASQLRLQNLRASCEALYSEMDPEERLR
FATTRHLRPMKTHLDRPLVVELGRDGARGPVGGKARPEAAEAPEGVDPPRRHHRHRD
KDKTPAAGDQDRAEAPKAESGEPGAREERPRPHRSKSKEAAGPPEARSERGRGPGPEG
GRRHHRGSPPEAAEREPRRHRAHRHQDPSKECAGAKGERRARHRGGPRAGPREAESG
EEPARRHRARHKAQPAHEAVEKETTEKEATEKEAEIVEADKEKELRNHQPREPHCDLE
TSGTVTVGPMHTLPSTCLQKVEEQPEDADNQRNVTRMGSSQPPDPNTIVHIPVMLTGPL
GEATVVPSPGNVDLESQAEGKKEVEADDVMRSGPRPIVPYSSMFCLSPNTLLRRFCHYI
VTMRYFEVVILVVIALSSIALAAEDPVRTDSPRNNAKYLDYIFTGVFTFEMVIK MID
LGLLLHPGAYFRDLWNILDFTVVSGALVAFASGSKGKDINTIKSLRVLRVLRPLKTI
KRLPKLKA VFDCVVNSLKNVNLIVYMLFMFIFAVIAVQLFKGKFFYCTDESKELER
DCRGQYLDYEKEEVEAQPRQWKKYDFHYDNVLWALLTLFTVSTGEGWPMVLKHSVDAT
YEEQGPSPGYRMELSIFYVVFVFPFFVNFVALIITFQEQQGDKVMSECSLEKNE
RACIDFAISAKPLTRYMPQNRQSFQYKTWTFVVSPPFEYFIMAMIALNTVVLMMKFYD
APYEYELMLKCLNIVFTSMFSMECVLKIIAFGVNLNYFRDAWNVDFVTVLGSITDILV
TEIAETNNFINLSFLRLFRAARLIKLLRQGYTIRILLWTFVQSFKALPYVCLLIAMLF
FTYAIIGMQVFGNIALDDDSINRHNNFRTFLQALMLLFRSATGEAWHEIMLSCLSNQ
ACDEQANATECGSDFAFYFVSFIFLCSFLMLNLFVAVIMDNFEYLTRDSSILGPHHL
DEFIRVWAEYDPAACGRISYNDFEMLKHMSPPLGLGKKCPARVAYKRLVRMNMPI SN
EDMTVHFTSTLMALIRTALEIKLAPAGTKQHQCDAELRKEISVVWANLPQKTLDLLVP
PHKPDEMTVGKVYAALMIFDFYKQNKTTTRDQMQQAPGGLSQMGPVSLFHPLKATLEQT
QPAVLRGARVFLRQKSSTSLNNGGAIQNQESGIKESVSWGTQRTQDAPHEARPPLERG
HSTEIPVGRSGALA VDVQMQSITRRGPDGEPQPGLESQGRAASMPRLAAETQPVTDAS
PMKRSISTLAQRPRGTHLCSTTPDRPPPSQASSHHHHHRCHRRRDRKQRSLEKGPSLS
ADMDGAPSSAVGPGLPPEGPTGCRERRERRQERGRSQERRQPSSSSSEKQRFYSCDR
FGGREPPKPKPSLSSHPTSPTAGQEPGHPQAGSAVGFPNTTPCCRETPSASPWPLAL
ELALTLTWGSVWTVRPLSTPCLRTRSLSRRLWPPTRAAPPGLPTCPP

FIGURE 22A

1 gatgtccga gctgtatcc ccggctcggc ccgggcagcc gccttctgag ccccgaccc
61 gaggcgccga gccgccgccg ccgatgggc tgggccgtgg agcgtctccg cagtcgtagc
121 tccagccgcc gcgcicccag ccccggcagc ctacagatca gcggcggcgg cgccggcggc
181 ggcgtctcc gcacgttcg ccgcagcgtg acccgagcc ctttctctt tgcagaatgg
241 cccgcttcgg agacgagatg ccggcccgt acgggggagg aggtccggg gcagccgccg
301 ggggtgtcgt gggcagcggg ggccggcgag gagccggggg cagccggcag ggccggcgagc
361 ccggggcgca aaggatgtac aagcagtcaa tggcgagag agcgcgacc atggcactct
421 acaaccccat ccccgctccg cagaactgcc tcacggttaa ccggtctctc ttcctctca
481 gcgaagacaa cgtggtgaga aaatacgcca aaaagatcac cgaatggcct cctttgaat
541 atatgattt agccaccatc atagcgaatt gcacgtctc cgcactggag cagcatctgc
601 ctgatgatga caagaccccg atgtctgaac ggctggatga cacagaacca tacttcattg
661 gaattttttg tticgaggct ggaattaaaa tcattgccct tgggtttgcc ttccacaaag
721 gctcctactt gaggaatggc tggaatgtca tggactttgt ggtggtgcta acgggcatct
781 tggcgacagt tgggacggag ttgacctac ggacgctgag ggcagttcga gtgctcgggc
841 cgctcaagct ggtgtctgga atcccaagtt tacaagctgt cctgaagtcg atcatgaagg
901 cgatgatccc ttgtctcgag atcgccctcc tctattttt tgcaatcctt atttttgcaa
961 tcatagggtt agaattttat atgggaaaat ttcataccac ctgctttgaa gaggggacag
1021 atgacatca ggggtgagtct ccggctccat gtgggacaga agagcccgcc cgcacctgcc
1081 ccaatgggac caaatgtcag cctactggg aaggggccaa caacgggac actcagttcg
1141 acaacatcct gtttgcagtg ctgactgtt tccagtgc ataatggaa ggggtgactg
1201 atctctcta caatagaac gatgcctcag ggaacacttg gaactggtt tacttcaccc
1261 cctcatcat catcggtccc tttttatgc tgaacctgt gctgggtgtg ctgtcagggg
1321 agtttgcga agaaaggga cgggtggaga accggcgggc ttttgaag ctgaggcggc
1381 aacaacagat tgaacgtgag ctcaatgggt acatggaatg gatctcaaaa gcagaagagg
1441 tgatcctcgc cgaggatgaa actgacgggg agcagaggca tccctttgat ggagctctgc
1501 ggagaaccac cataaagaaa agcaagacag atttgctcaa cccgaagag gctgaggatc
1561 agctggctga tatagcctt gtgggttctc ccttcgccc agccagcatt aaaagtgcc
1621 agctggagaa ctgcacctt ttacacaaa aggagaggag gatcggttc tacatccgcc
1681 gcattgtcaa aactcaggcc ttctactgga ctgtactcag ttggtagct ctcaacacgc
1741 tgtgtgtgc tattgttac tacaaccagc ccgagtggct ctccgacttc ctttactatg
1801 cagaattcat ttcttagga ctctttatgt ccgaaatgt tataaaaatg tacgggcttg
1861 ggacgggcc ttactccac tcttctca actgcttga ctgtggggtt atcattggga
1921 gcattctga ggtcatctgg gctgtcataa aacctggcac atcctttgga atcagcgtgt
1981 tacgagccct cagggtattg cgtatttca aagtcacaaa gtactgggca tcttcagaa
2041 acctggtcgt ctctctcctc aactccatga agtccatcat cagcctgttg ttctcctt
2101 tctgttcat tgcgtcttc gcccttttg gaatgcaact ctccggcggc cagttaatt
2161 tcatgaagg gactcctccc accaacttcg atactttcc agcagcaata atgacggtgt
2221 ttcagatcct gacggcgga gactggaac aggtcatgta cgacgggac aagtctcagg
2281 gggcggtgca gggcggcag gtgtctcca tctatttcat tgtactgac ctctttggga
2341 actacacct cctgaatgt ttcttgcca tctgtgtga caatctggcc aacggccagg
2401 agctcacaa ggtggaggcg gacgagcaag aggaagaaga agcagcgaac cagaaacttg
2461 ccctacagaa agccaaggag gtggcagaag tgaatcctt gtccgcgcc aacatgtcta
2521 tagctgtgaa agagcaacag aagaatcaa agccagccaa gtccgtgtgg gagcagcgga
2581 ccagtगत gcgaaagcag aactgtctgg ccagccggga ggccctgtat aacgaaatgg
2641 acccgagca gcgtggaag gctgcctaca cgcggcacct gcggccagac atgaagacgc

FIGURE 22B

2701 acttggaccg gccgctggtg gtggaccgc aggagaaccg caacaacaac accaacaaga
2761 gccgggaggc cgagcccacc gtggaccagc gcctcggcca gcagcgcgc gaggacttcc
2821 tcaggaaaca ggcccgtac cagatcggg cccgggacc cagcggctcg gcgggcctgg
2881 acgcacggag gccctgggag ggaagccagg aggccgagct gagccgggag ggaccctacg
2941 gccgcgagtc ggaccaccac gcccgggagg gcagcctgga gcaacccggg ttctgggagg
3001 gcgaggccga gcgaggcaag gccggggacc cccaccggag gcacgtgcac cggcaggggg
3061 gcagcaggga gaggcgagc ggggtcccgc gcacgggggc ggacggggag catcgacgtc
3121 atcgccgcga ccgcaggccc ggggaggagg gtccggaggga caaggcggag cggagggggc
3181 ggaccgcga gggcagccgg ccggccggg gcggcgaggg cgaggggag ggccccgacg
3241 ggggcgagcg caggagaagg caccggcatg gcgtccagc cagtagag ggaggacgcgc
3301 ggaggaggga caaggagcgg aggcacgga ggaggaaaga gaaccagggc tccgggggtcc
3361 ctgtgtcggg ccccaacctg tcaaccacc ggccaatcca gcaggacctg ggccgccaag
3421 acccaccctt ggagaggat attgacaaca tgaagaaca caagctggcc accgcggagt
3481 cggccgtcc ccacggcagc ctggccacg ccggcctgcc ccagagcca gccaagatgg
3541 gaaacagcac cgacccggc ccatgttg ccacccctgc catggccacc aacccccaga
3601 acgccgccag ccggcgagc ccaacaacc cggggaacc atccaatcc ggcccccca
3661 agacccccga gaatagctt atcgtacca acccagcgg caccagacc aattcagcta
3721 agactgccag gaaacccgac cacaccagc tggacatcc cccagcctgc cccccccc
3781 tcaaccacac cgtcgtacaa gtgaacaaa acgccaacc agaccactg ccaaaaaag
3841 aggaagagaa gaaggaggag gaggaagac accgtggga agacggcct aagccaatgc
3901 ctccctatag ctccatgtt atcctgtcca cgaccaacc ccttcggc ctgtgccatt
3961 acatctgaa cctgcgtac ttgagatgt gcacctcat ggtcattgcc atgagcagca
4021 tcgcccaggc cgccgaggac cctgtgcag ccaacgcacc tcggaacaac gtgtgcgat
4081 actttgacta cgtttttaca ggcgtctca cctttgagat ggtgatcaag atgattgacc
4141 tggggtcgt cctgcatcag ggtgcctact tccgtgacct ctggaatatt ctgcattca
4201 tagtggtagc tggggccctg gtagccttg ccttacttg caatagcaaa ggaagagaca
4261 tcaacacgat taaatccctc cgagtcctcc ggggtctacg acctctaaa accatcaagc
4321 ggtgccaata gctcaaggct gtgttgact gtgtggtgaa ctacttaaa aacgtctca
4381 acatctcat cgtctacatg ctattcatgt tcatctcgc cgtgtgtgct gtgcagctt
4441 tcaaggggaa attcttcac tgcactgac agtccaaaga gtttgagaaa gattgtcgag
4501 gcaaatacct cctctacgag aagaatgagg tgaaggcgc agaccgggag tgaagaagt
4561 atgaattcca ttacgacaat gtgtgtggg ctctgtgac cctctcacc gtgtccacgg
4621 gagaaggctg gccacagtc ctcaagcatt cgggtggagc cacctttgag aaccagggcc
4681 ccagccccgg gtaccgatg gagatgtcca tttctacgt cgtctactt gtgtgttcc
4741 ccttctctt tgtcaatac ttgtggcct tgatcatcat cacctccag gagcaagggg
4801 acaagatgat ggaggaatac agcctggaga aaaatgagag ggcctgcatt gattcgcca
4861 tcagcgccaa gccgtgacc cgacacatgc cgcagaaca gcagagctt cagtaccgca
4921 tgtggcagtt cgtggtgtc ccgcttctg agtacagat catggccatg atcgccctca
4981 acaccatcgt gcttatgatg aagtctatg gggcttctgt tgcttatgaa aatgccctgc
5041 ggggtgtcaa catcgtctc acctccctct tctcttgga atgtgtgtg aaagtcatgg
5101 cttttggat tctgaatat tccgcgatg cctggaacat ctctgactt gtgactgtc
5161 tgggcagcat caccgatatc ctctgactg agtttggga tccgaataac tcatcaacc
5221 tgagctttc ccgctctc cgagctgcc ggctcatcaa acttctccg cagggttaca
5281 ccatccgat tcttctctg acctttgtg agtcttcaa ggccctgcct tatgtctgc
5341 tctgtatgc catgctctt tcatctatg ccatcattg gatgcagggtg tttgtaaca

FIG. 22C

5401 ttggcatcga cgtggaggac gaggacagt atgaagatga gttccaaatc actgagcaca
5461 ataatctccg gacctcttc caggccctca tgcttctt ccggagtgcc accggggaag
5521 cttggcacia catcatgctt tctgcctca gcgggaacc gtgtgataag aactctggca
5581 tctgactcg agagtgtggc aatgaattg cttatttta cttgtttcc ttcatttcc
5641 tctgctcgtt tctgatgtg aatctcttg tcgccgtcat catggacaac tttagtacc
5701 tcaccgaga ctctccatc ctgggcccc accacctgga tgagtacgtg cgtgtctggg
5761 ccgagtatga ccccgagct tggggccgca tgccttacct ggacatgat cagatgctga
5821 gacacatgct tccgcccctg ggtctgggga agaagtgtcc ggccagagtg gcttacaagc
5881 ggcttctgcg gatggacctg cccgtcgag atgacaacac cgtccacttc aattccacc
5941 tcatggctct gatccgcaca gccctggaca tcaagattgc caaggagga gccgacaaac
6001 agcagatgga cgctgagctg cgggaaggaga tgatggcgat ttggcccaat ctgtccaga
6061 agacgctaga cctgtgtg acacccaca agtcacgga cctaccgtg gggaagatct
6121 acgagccat gatgatcatg gactactacc ggagagcaa ggccaagaag ctgaggcca
6181 tgcgcgagga gcaggaccgg acaccctca tgtccagcg catggagccc ccgtcccaa
6241 cgcaggaagg gggacctggc cagaacgcc tccctccac ccagctggac ccaggaggag
6301 cctgatggc tcagaaagc ggctcaagg agagcccgtc ctgggtgacc cagcgtgcc
6361 aggagatgtt ccagaagacg ggcacatgga gtccggaaca agggccccct accgacatgc
6421 ccaacagcca gcctaactct cagtccgtg agatcgaga gatgggcaga gatggctact
6481 ccgacagcga gcaactct cccatggaag gccaggggcg ggctgcctcc atccccgcc
6541 tccctgcaga gaaccagagg agaaggggcc ggccacgtgg gaataacctc agtaccatct
6601 cagacaccag cccatgaag cgttcagct ccgtgctgg cccaaggcc cgagcctgg
6661 acgattactc gctggagcgg gtccgccc aggagaacca gggcaccac cagcggcgcc
6721 gcgaccgag ccaccgccc tctgagcgt cctggggcg ctacaccgat gtgacacag
6781 gcttggggac agacctgagc atgaccacc aatccgggga cctgccgtc aaggagcggg
6841 accaggagcg gggccggccc aagatcgga agcatcgaca gcaccaccac caccaccac
6901 accaccacca tccccgccc cccgacaagg accgctatgc ccaggaacgg cgggaccag
6961 gccgggcacg ggctcgggac cagcgtggt cccgtcgcc cagcgaggcg cgagagcaca
7021 tggcgaccg gcagggcagt agttcgtaa gtggaagccc agccccctca acatctggta
7081 ccagcactcc gcggcggggc cgccgccagc tccccagac ccctccacc cccggccac
7141 acgtgtccta tccccgtg atccgtaagg ccggcggctc ggggcccccg cagcagcagc
7201 agcagcagca gcagcagcag caggcgggtg ccaggccggg ccggcgggcc accagcggcc
7261 ctggaggta cccaggcccc acggccgagc ctctggccgg agatcgggcg cccacggggg
7321 gccacagcag cggccgctc cccaggatgg agaggcgggt cccaggcccc gcccggagcg
7381 agtccccag ggctgtcga caggcgggg cccgtggcc ggcatctggc ccgacgtgt
7441 ccgaggggcc cccgggtccc cggcaccatg gctactacc gggctccgac tacgacgag
7501 ccgatggccc gggcagcggg ggccgagag aggccatggc cggggcctac gacgcgccac
7561 cccccgtac acacgcgtcc tcggcgcca ccggcgctc gccaggact cccgggcct
7621 cgggccccgg ctgcgctc cttctcggc acggccggcg actcccaac ggctactacc
7681 cggcgacagg actggccagg cccgcgggc cgggtccag gaagggcctg cacgaacctt
7741 acagcgagag tgacgatgat tgggtctaag cccggcgag gtggcgccc cccggcccc
7801 cagcacc

FIGURE 22D

MARFGDEMPARYGGGGSGAAAGVVVGSGGGRGAGGSRQGGQPGA
QRMYSQSMARARTMALYNPIPVQRNCLTVNRSFLFSEDNVVRKYAKKITEWPPFEY
MILATIIANCIVLALEQHLRDDDKTPMSERLDDTEPYFIGIFCFEAGIKIILGFAFH
KGSYLRNGWNVMDFVVVLTGILATVGTEFDLRTLRAVRVLRPLKLVSGIPSLQVVLKS
IMKAMPLLQIGLLLFFAILIFAIIGLEFYMGKFHTTCFEEGTDDIQGESAPCGTEE
PARTCPNGTKCQPYWEGPNNGITQFDNLFAVLTVFQCITMEGWTDLLYNSNDASGNT
WNWLYFIPLIIIGSFFMLNLVLGVLSGEFAKERERVENRRAFLKLRRQQQIERELNGY
MEWISKAEEVILAEDETDGEQRHPFDGALRRTTIKKSKTDLLNPEEAEDQLADIASVG
SPFARASIKSAKLENSTFFHKKERRMRFYIRRMVKTQAFYWTVLSLVALNTLCVAIVH
YNQPEWLSDFLYYAEFIFLGLFMSEMFIMYGLGTRPYFHSSFNCFDCGVIIGSIFEV
IWAVIKPGTSFGISVLRALRLLRIFKVTKYWASLRNLVVSLLNSMKSIISLLFLLFLF
IVVFALLGMQLFGGQFNFDGTPPTNFDTFPAAIMTVFQILTGEDWNEVMYDGKSQG
GVQGGMVFSIYFIVLTLFGNYTLLNVFLAIAVDNLANAQELTKVEADEQEEEEAAANQK
LALQAKEVAEVSPLSAANMSIAVKEQQKNQKPAKSVWEQRTSEMRKQNLASREALY
NEMDPDERWKAAYTRHLRPDMKTHLDRPLVVDPOENRNNNTNKSRAAEPTVDQRLGQQ
RAEDFLRKQARYHDRARDPSGSAGLDARRPWAGSQEAELSREGPYGRESDDHAREGSL
EQPGFWEGEAERGKAGDPHRRHVHRQGGSRSSRSGSPRTGADGEHRRHRAHRRPGEEG
PEDKAERRARHREGSRPARGGEGEGEPDGGERRRRHRHGAPATYEGDARREDKERRH
RRRKENQGGSGVPVSGPNLSTTRPIQQDLGRQDPPLAEDIDNMKNKLATAESAAPHGS
LGHAGLPQSPAKMGNSTDPGPMALPAMATNPQNAASRRTPNPNPGNPSNPGPPKTPEN
SLIVTNPSGTQTNSAKTARKPDHTTVDIPPACPPPLNHTVVQVNKNANPDPLPKKEEE
KKEEEEDDRGEDGPKPMPPYSSMFLSTTNPLRRLCHYLNLRYFEMCILMVIAMSSI
ALAAEDPVQPNAPRNNVLR YFDYVFTGVFTFEMVIKMIDLGLVLHQGA YFRDLWNILD
FTVVSGALVAFVFTGNSKGDINTIKSLRVLRLPLKTIKRLPKLKA VFDCVVNSLK
NVFNILIVYMLFMFIFAVVA VQLFKGKFFHCTDESKEFEKDCRGKYLLYEKNEVKARD
REWKKYEFHYDNVLWALLTLFTVSTGEGWPQVLKHSVDATFENQGPSPGYRMEMSIFY
VVYFVVPFFFFVNI FVALIITFQE QGDKMMEEYSLEKNERACIDFAISAKPLTRHMP
QNKQSFQYRMWQFVVSPPFEYTIMAMIALNTIVLMMKFY GASVAYENALRVFNTVFTS
LFSLECVLKVMAFGILNYFRDAWNIFDFVTVLGSITDILVTEFGNPNNFNLSFLRLF
RAARLIKLLRQGYTRILLWTFVQSFKALPYVCLLIAMLFFIYAIIGMQVFIGNIDV
EDEDSDDEFQITEHNNFR TFFQALMLLFRSATGEA WHNIMLSCLSGKPCDKNSGILT
RECGNEFA YFYFVSFIFLCSFLMLNLFVAVIMDNFEYLTRDSSILGPHHLDEYVRVWA
EYDPAAWGRMPYLDMYQMLRHMSPPLGLGKKCPARVAYKRLLRMDLPVADDNTVHFNS
TLMALIRTALDIKIAKGGADKQQMDAELRKEMMAIWPNL SQKTLDLLVTPHKSTDLT
GKIYAAMMIMEYYRQSKAKKLQAMREEQDRTPLMFQRMPPSPTQEGGPGQNALPSTQ
LDPGGALMAHESGLKESPSWVTQRAQEMFQKTGTWSPEQGPPTDMPNSQPNQSQSVEMR
EMGRDGYSDSEHYLPMEGQGRAASMPRLPAENQRRRGRPRGNNLSTISDTSPMKRSAS
VLGPKARRLDDYSLERVPPEENQRHHQRRRDRSHRASERSLGRYTDVDTGLGTDLSMT
TQSGDLPSKERDQERGRPKDRKHRQH HHHHHHHHHPPPPDKDRYAQERPDHGRARARD
QRWSRSPSEGREHMAHRQGGSSVS GSPAPSTSGTSTPRRGRRQLPQTPSTPRPHVSYS
PVIRKAGGSGPPQQQQQQQQQQA VARPGRAATSGPRRYPGPTAEPLAGDRPPTGGHS
SGRSPRMERRVPGPARSESPRACRHGGARWPASGPHVSEGPPGPRHHGYRGS DYDEA
DGP GSGGEEAMAGAYDAPPPVRHASSGATGRSPRTPRASGPACASPSRHGRRLPNGY
YPAHGLARPRGPGSRKGLHEPYSESDDDDWC

FIGURE 23A

1 gatgtccga gctgctatcc ccggctcggc ccgggcagcc gccttctgag cccccgaccc
61 gaggcgccga gccgccgccg ccgcatgggc tgggccgtgg agcgtctccg cagtcgtagc
121 tccagccgcc gcgctcccag ccccggcagc ctacagatca gcggcggcgg cgccggcggc
181 ggctgttcc gcatcgttcg ccgcagcgtc accggagacc cttgtcttt tgcagaatgg
241 ccgcttcgg agacgagatg ccggcccgtc acgggggagg aggtccggg gcagccgccg
301 ggggtggctgt gggcagcggg ggccggcgag gagccggggg cagccggcag ggccggcagg
361 ccggggcgca aaggatgtac aagcagtcaa tggcgagag agcggcgacc atggcactct
421 acaaccccat ccccgctcca cagaactgcc tcacggtaa ccggctctc ttcctttca
481 gcgaagacaa cgtggtgaga aaatacgcca aaaagatcac cgaatggcct cctttgaat
541 atatgattt agccaccatc atagcgaatt gcatcgtcct cgcactggag cagcatctgc
601 ctgatgatga caagaccccg atgtctgaac ggctggatga cacagaacca tacttcattg
661 gaatttttg ttcgaggct ggaataaaa tcattgccct tgggtttgcc ttccaaaag
721 gctcctactt gaggaatggc tggaatgta tggacttgt ggtggtgcta acgggcatct
781 tggcgacagt tgggacggag ttgacctac ggacgtgag ggacgttga gtgctgcggc
841 cgtcaagct ggtgtctgga atcccaagt tacaagctc cctgaagtc atcatgaagg
901 cgatgatccc ttgtgcag atcgccctcc tctatttt tgcaatcct attttgcga
961 tcatagggtt agaattttat atgggaaaat tcataccac ctgcttgaa gaggggacag
1021 atgacattca ggtgagctc ccggctccat gtgggacaga agagccgcc cgcacctgcc
1081 ccaatgggac caaatgtcag ccctactggg aagggcccaa caacgggac actcagtgc
1141 acaacatcct gttgcagtg ctgactgtt tccagtgc atccatggaa ggtggtgactg
1201 atcctctca caatgaac gatgcctcag ggaacactg gaactggtg tacttcaccc
1261 cctcatcat catcggtccc tttttatgc tgaacctgt gctgggtgtg ctgtcgggg
1321 agtttgcga agaaaggga cgggtggaga accggcgggc tttctgaag ctgaggcggc
1381 aacaacagat tgaactgtg ctcaatgggt acatggaat gatctaaaa gcagaagagg
1441 tgatcctgc cgaggatgaa actgacgggg agcagaggca tccctttgat ggagctctgc
1501 ggagaaccac cataaagaaa agcaagacag atttgtcaa ccccgagag gctgaggatc
1561 agctggctga tatagcctc gtgggtctc ccttcggccg agccagcatt aaaagtcca
1621 agctggagaa ctgcacctt tttcaaaaa aggagaggag gatcgcttc tacatccgcc
1681 gcatggtaaa aactcaggcc ttctactgga ctgtactcag ttggtagct ctcaacacgc
1741 tgtgtgttc tattgttac tacaaccagc ccgagtggct ctccgacttc cttactatg
1801 cagaattcat ttttagga ctctttatg ccgaatgtt tataaaaatg tacgggcttg
1861 ggacgcggcc ttactccac tcttcttca actgcttga ctgtggggt atcattggga
1921 gcatcttga ggcatctgg gctgtcataa aacctggcac atcctttgga atcagcgtgt
1981 tacgagccct caggttattg cgtatttca aagtcacaaa gtactgggca tctctcagaa
2041 acctggtcgt ctctctctc aactccatga agtccatcat cagcctgtt tttctcttt
2101 tctgttcat tctgtcttc gcccttttg gaatgcaact ctccggcggc cagtttaatt
2161 tcgatgaagg gactcctccc accaacttcg atactttcc agcagcaata atgacggtgt
2221 ttcagatcct gacggcgga gactggaacg aggtcatgta cgacgggac aagtctcagg
2281 ggggcgtgca gggcgcatg gtgttctcca tctattcat tgtactgac ctctttggga
2341 actacacct cctgaatgtg ttcttgcca tcgctgtgga caatctggcc aacgccagg
2401 agtcaccaa ggtggaggcg gacgagcaag aggaagaaga agcagcgaac cagaaacttg
2461 ccctacagaa agccaaggag gtggcagaag tgagtctct gtccgcgcc aacatgtcta
2521 tagctgtgaa agagcaacag aagaalcaaa agccagcaa gtccgtgtg gagcagcggg
2581 ccagttagat gcgaaagcag aactgtctg ccagccggga ggccctgtat aacgaaatgg
2641 acccgagcga gcgttggaag gctgcctaca cgcggcacct gcggccagac atgaagacgc
2701 acttgaccg gccgctggtg gtggaccgc aggagaaccg caacaacaac accaacaaga

FIGURE 23B

2761 gccgggcggc cgagcccacc gtggaccagc gcctcggcca gcagcgccc gaggacttc
 2821 tcaggaaaca ggcccgtac cacgatcggg cccgggaccc cagcggctcg gcgggcctgg
 2881 acgcacggag gccctggcg ggaagccagg aggcgagct gagccgggag ggaccctacg
 2941 gccgcgagtc ggaccaccac gcccgggagg gcagcctgga gcaaccggg ttctgggagg
 3001 gcgaggccga gcgaggcaag gccggggacc cccaccggag gcacgtgcac cggcaggggg
 3061 gcagcaggga gagccgcagc gggtccccgc gcacgggcgc ggacggggag calcgacgtc
 3121 atcgcgcgca ccgcaggccc ggggaggagg gtccggaggga caaggcggag cggagggcgc
 3181 ggaccgcga gggcagccgg ccggcccggg gcggcgaggg cgagggcgag ggccccgacg
 3241 ggggcgagcg caggagaagg caccggcatg gcgtccagc cacgtacgag ggggacgcgc
 3301 ggaggaggga caaggagcgg aggcacgga ggaggaaaga gaaccagggc tccggggtcc
 3361 ctgtgtcggg ccccaacctg tcaaccacc ggccaatcca gcaggacctg ggccgccaag
 3421 acccaccctt ggagaggat attgacaaca tgaagaaca caagctggcc accgcggagt
 3481 cggccgtcc ccacggcagc ctgtgccag ccggcctgcc ccagagccca gccaatgg
 3541 gaaacagcac cgacccggc ccatgtctg ccatccctgc catggccacc aacccccaga
 3601 acgccgcag ccgccggacg ccaacaacc cggggaacc atccaatccc ggcccccca
 3661 agacccccga gaatagcctt atgtcacca acccagcgg caccagacc aattcagcta
 3721 agactgccag gaaaccgac cacaccacag tggacatccc cccagcctgc cccccccc
 3781 tcaaccacac cgtctgataa gtgaacaaaa acgccaacc agaccactg ccaaaaaag
 3841 aggaagagaa gaaggaggag gaggaagacg accgtgggga agacggcct aagccaatg
 3901 ctccctatag ctccatgtt atcctgtcca cgaccaacc ccttcgccg ctgtgccatt
 3961 acatctgaa cctgcgtac ttgagatgt gcacctcat ggtcattgcc atgagcagca
 4021 tcgccctggc cgccgaggac cctgtgcag ccaacgcacc tcggaacaac gtgtgcgat
 4081 actttgacta cgttttaca ggcgtctca ctttgagat ggtgatcaag atgattgacc
 4141 tggggctcgt cctgcatcag ggtgcctact tccgtgacct ctggaatatt ctgacttca
 4201 tagtggtag tggggccctg gtagccttg ccttacttg caatagcaa ggaaaagaca
 4261 tcaacacgat taaatccctc cgagtctcc ggggtctacg acctttaa accatcaagc
 4321 gggtgcaaaa gctcaaggct gtgttgact gtgtggtgaa ctacttaa aacgtctca
 4381 acatctcat cgtctacatg ctattcatgt tcatcttcgc cgtggtggct gtgcagctct
 4441 tcaaggggaa attcttcac tgcactgacg agtccaaaga gtttgagaaa gattgtcgag
 4501 gcaaatacct cctctacgag aagaatgagg tgaaggcgcg agaccgggag tggagaagt
 4561 atgaattcca ttacgacaat gtgtgtggg ctctgtgac cctcttacc gtgtccacgg
 4621 gagaaggctg gccacagtc ctcaagcatt cgggtggacg caccttgag aaccagggcc
 4681 ccagccccgg gtaccgatg gagatgtcca tttctacgt cgtctactt gtggtgttc
 4741 ccttctctt tgtcaatac ttttggcct tgatcatcat caccttcag gagcaagggg
 4801 acaagatgat ggaggaatac agcctggaga aaaatgagag ggcctgcatt gatttcgca
 4861 tcagcgccaa gccgtgacc cgacacatgc cgcagaaca gcagagcttc cagtaccgca
 4921 tgtggcagtt cgtggtgtct ccgccttcg agtacagat catggccatg atcgccctca
 4981 acaccatcgt gcttatgatg aagttctatg gggcttctgt tgcattgaa aatgccctgc
 5041 ggtgttcaa catgtcttc acctccctct tctcttgga atgtgtgtg aaagtcattg
 5101 cttttggat tctgaattat ttccgcatg cctggaacat ctctgactt gtgactgtc
 5161 tgggcagcat caccgatc ctctgactg agtttgggaa tccgaataac tcatcaacc
 5221 tgagctttct ccgctcttc cgagtgccc ggctcatcaa acttctcgt cagggttaca
 5281 ccatccgat tcttcttg acccttgc agtccitcaa ggccctgcct tatgtctgc
 5341 tctgatcgc catgtcttc tcatctatg ccatcattgg gatgcagggtg ttgtgaaca

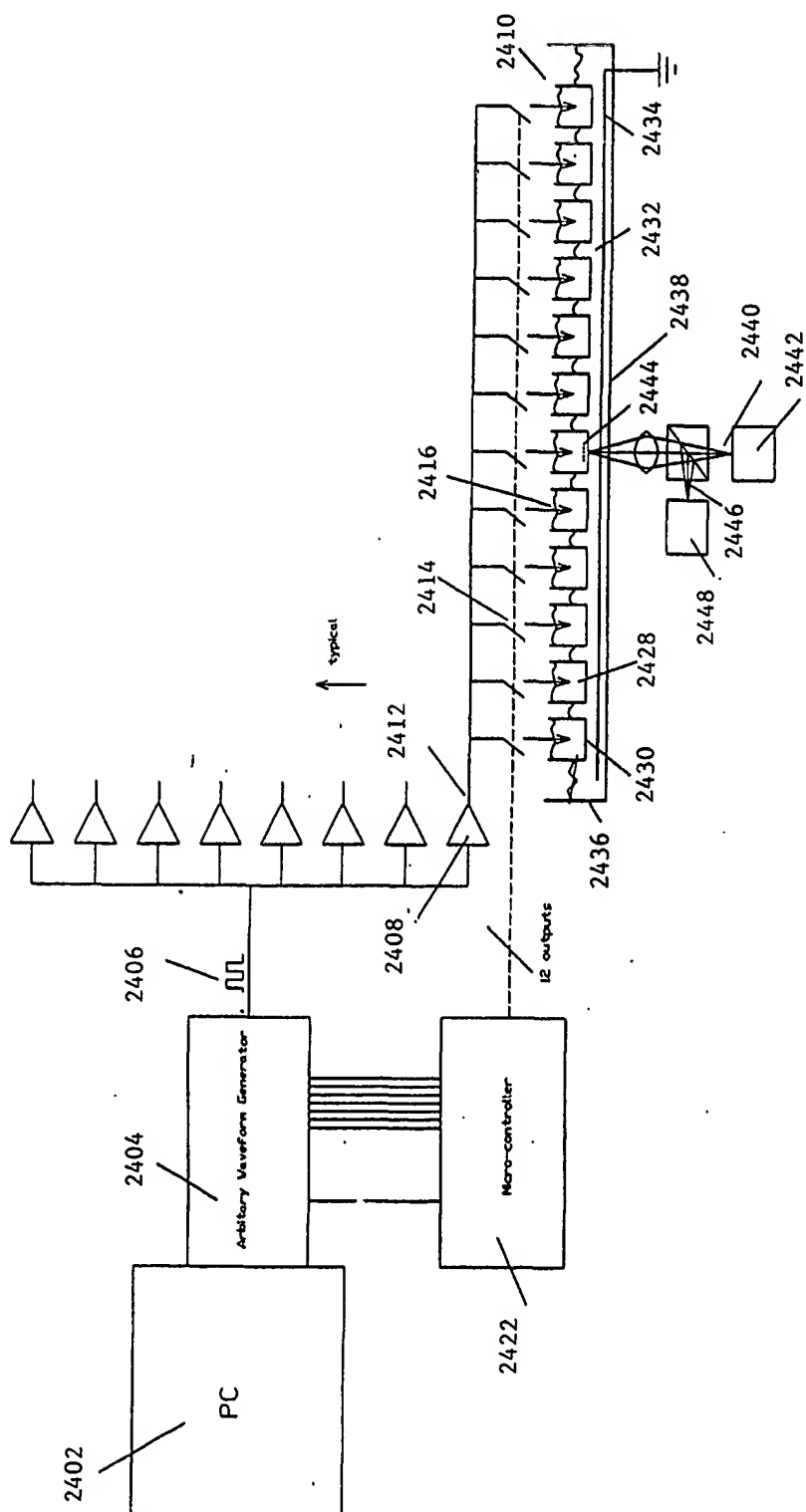
FIG. 23C

5401 ttggcatcga cgtggaggac gaggacagt atgaagatga gtccaaatc actgagcaca
 5461 ataacttccg gaccttctc caggccctca tgcttctctt ccggagtgcc accggggaaag
 5521 ctgggcacaa catcatgctt tctgacctca gcgggaaacc gtgtgataag aactctggca
 5581 tctgactcg agagtgtggc aatgaattg cttatttta cttgtttcc ttcattctcc
 5641 tctgctcgtt tctgatgctg aatctctttg tcgccgtcat catggacaac tttagtacc
 5701 tcacccgaga ctctccatc ctgggcccc accacctgga tgagtacgtg cgtgtctggg
 5761 ccgagtatga ccccgagct tggggccgca tgccttacct ggacatgat cagatgctga
 5821 gacacatgtc tccgccccg ggtctgggga agaagtgtcc ggccagagtg gcttacaagc
 5881 ggcttctcg gatggacctg ccgctgcag atgacaacac cgtccacttc aattccacc
 5941 tcatggctct gatccgcaca gccctggaca tcaagattgc caaggaggga gccgacaaac
 6001 agcagatgga cgtgagctg cggaaggaga tgatggcgat ttggcccaat ctgtccaga
 6061 agacgctaga cctgctgtc acacctaca agtcacgga cctcaccgtg gggaagatct
 6121 acgcagccat gatgatcatg gactactacc ggacagagca ggccaagaag ctgcaggcca
 6181 tgcgcgagga gcaggaccgg acaccttca tgtccagcg catggagccc ccgtcccaa
 6241 cgagggaagg gggacctggc cagaacgccc tccccccac ccagctggac ccaggaggag
 6301 ccctgatggc tcacgaaagc ggctcaagg agagcccgct ctgggtgacc cagcgtgccc
 6361 aggagatgtt ccagaagacg ggcacatgga gtccggaaca agggccccct accgacatgc
 6421 ccaacagcca gcctaactc cagtccgtgg agatcgaga gatgggcaga gatggctact
 6481 ccgacagcga gcactacct cccatggaag gccaggggcg ggctgcctcc atgccccgcc
 6541 tcctgcaga gaaccagagg agaaggggcc ggccacgtgg gaataacctc agtaccatct
 6601 cagacaccag ccccatgaag cgttcagct ccgtgctggg cccaaggcc cgacgcctgg
 6661 acgattactc gctggagcgg gtccccccc aggagaacca gcggcaccac cagcggcgcc
 6721 gcgaccgag ccaccgcgc tctgagcgt ccctggggcg ctacaccgat gtggacacag
 6781 gcttggggac agacctgagc atgaccacc aatccgggga cctgccgtcg aaggagcggg
 6841 accaggagcg gggccggccc aaggatcgga agcatcgaca gcaccaccac caccaccac
 6901 accaccacca tccccgcc cccgacaagg accgctatgc ccaggaacgg ccggaccacg
 6961 gccgggcacg ggctcgggac cagcgtggt cccgctgcc cagcaggggc cgagagcaca
 7021 tggcgaccg gcagtagtic cgtaatgga agccagccc cctcaacatc tggtagcagc
 7081 actccgccc gggggcccg ccagctccc cagacccct ccacccccg gccacacgtg
 7141 tctattccc ctgtgatcc taaggccggc ggctcggggc cccgcagca gcagcagcag
 7201 cagcagcgcg tggccaggcc gggccggcg gccaccagcg gccctcggag gtaccaggcc
 7261 cccacggcg agcctctggc cggagatcgg ccgcccacgg ggggccacag cagcggccgc
 7321 tcgccagga tggagaggcg ggtcccagg ccggcccga gcgagtccc cagggcctgt
 7381 cgacacggcg gggcccggg gccggcatct ggcccgcag gtccgaggg gccccgggt
 7441 ccccggcacc atggctacta ccggggctcc gactacgac aggccgatgg cccgggcagc
 7501 gggggcgcg aggaggccat ggccggggc tacgacgcg cccccccgt acgacagcg
 7561 tctcggggc ccaccggcg ctgcccagg actccccgg cctcggggc ggctgcgcc
 7621 tcgcttctc ggcacggcg gcgactccc aacggctact accggcgca cggactggcc
 7681 agggcccgcg ggcggggctc cagggaaggc ctgcacgaac cctacagca gagtgcgat
 7741 gattgtgct aagccgggc gaggtggcg ccggccggc cccacgcac c

FIG. 23D

MARFGDEMPARYGGGGSGAAAGVVVVGSGGGRGAGGSRQGGQPGA
QRMYSMAQRARTMALYNPIVRQNCLTVNRSFLFSEDNVVRKYAKKITEWPPFEY
MLATIIANCIVLALEQHLRDDDKTPMSERLDDTEPYFIGIFCFEAGIKIILGFAFH
KGSYLRNGWNVMDFFVVLGTGILATVGTEFDLRTLRAVRVLRPLKLVSGIPSLQVVLKS
IMKAMPLLQIGLLLFFAILFAIIGLEFYMGKFHTTCFEEGTDDIQGESAPCGTEE
PARTCPNGTKCQPYWEGPNNGITQFDNLFAVLTVFQCITMEGWTDLLYNSNDASGNT
WNWLYFIPLIIIGSFFMLNLVLGVLSGEFAKERERVENRRAFLKLRRQQQIERELNGY
MEWISKAEEVILAEDETDGEQRHPFDGALRRTTIKKSKTDLLNPEEAEDQLADIASVG
SPFARASIKSAKLENSTFFHKKERRMRFYRRMVKTQAFYWTVLSLVALNTLCVAIVH
YNQPEWLSDFLYYAEFIFLGLFMSEMFIMYGLGTRPYFHSSFNCFDCGVIIGSIFEV
IWAVIKPGTSFGISVLRALRLRIFKVTKYWASLRNLVVSLLNSMKSIISLLFLLFLF
IVVFALLGMQLFGGQFNDEGTPPTNFDTFPAAIMTVFQILTGEDWNEVMYDGIKSQG
GVQGGMVFSIYFVLTFLGNYTLLNVFLAIAVDNLANAQELTKVEADEQEEEEEAANQK
LALQKAKEVAEVSPLSAANMSIAVKEQQKNQKPAKSVWEQRTSEMRKQNLLASREALY
NEMDPDERWKAAYTRHLRPDMKTHLDRPLVVDPQENRNNNTNKSRAAEPTVDQRLGQQ
RAEDFLRKQARYHDRARDPSGSAGLDARRPWAGSQEAELSREGPYGRESDDHAREGSL
EQPGFWEGEAERGKAGDPHRRHVHRQGGSRRESRSGSPRTGADGEHRRHRAHRRPGEEG
PEDKAERRARHREGSRPARGEGEGEGPDGGERRRRHRHGAPATYEGDARREDKERRH
RRRKENQGSQVVPVSGPNLSTTRPIQQDLGRQDPPLAEDIDNMKNKLATAESAAPHGS
LGHAGLPQSPAKMGNSTDPGPMALPAMATNPQNAASRRTPNPNPGNPSNPGPKTPEN
SLIVTNPSGTQTNNAKTARKPDHTTVDIPACPPPLNHTVVQVNKNANPDPLPKKEEE
KKEEEEDDRGEDGPKPMPPYSSMFLSTTNPLRLCHYLNLRYFEMCILMVIAMSSI
ALAAEDPVQPNAPRNNVLRVFDYVFTGVFTFEMVIKMDLGLVLHQGAYFRDLWNILD
FTVVSAGALVAFATGNSKKGKDINTIKSLRVLRVLRPLKTIKRLPKLKA VFDCVVSNLK
NVFNILIVYMLFMFIFAVVA VQLFKGKFFHCTDESKEFEKDCRGKYLLYEKNEVKARD
REWKKYEFHYDNVLWALLTLFTVSTGEGWPQVLKHSVDATFENQGPSPGYRMEMSIFY
VVYFVVPFFVNFVALIITFQEQGDKMMEEYSLEKNERACIDFAISAKPLTRHMP
QNKQSFQYRMWQFVVSPPFEYTIMAMIALNTIVLMMKFYGASVAYENALRVFNIVFTS
LFSLECVLKVMAFGILNYFRDAWNIFDFVTVLGSITDILVTEFGNPNPNFINLSFLRLF
RAARLIKLLRQGYTIRILLWTFVQSFKALPYVCLLIAMLFFTYAIGMQVFGNIGIDV
EDEDSEDEFQITEHNNFRFTFFQALMILLFRSATGEA WHNIMLSCLSGKPCDKNSGILT
RECGNEFAYFYFVSFIFLCSFLMLNLFAVIMDNFEYLTRDSSILGPHHLDEYVRVWA
EYDPAAWGRMPYLDMYQMLRHMSPPGLGKKCPARVAYKRLLRMDLPVADDNTVHFNS
TLMALIRALTALDIKAKGGADKQQMDAELRKEMMAIWPNLQKTLDDLVTPHKSTDLTV
GKIYAAMMIMEYYRQSKAKKLQAMREEQDRTPLMFQRMPEPSPTQEGGPGQNALPSTQ
LDPGGALMAHESGLKESPSWVTQRAQEMFQKTGTWSPEQGPPTDMPNSQPNQSQSVEMR
EMGRDGYSDSEHYLPMEGQGRAASMPRLPAENQRRRGRPRGNLSTISDTSPMKRSAS
VLGPKARRLDDYSLERVPPEENQRHHQRRRDRSHRASERSLGRYTDVDTGLGTDLSMT
TQSGDLPSKERDQERGRPKDRKHRQHHHHHHHHHHPPPDKDRYAQERPDHGRARARD
QRWSRSPSEGREHMAHRQ

FIG. 24



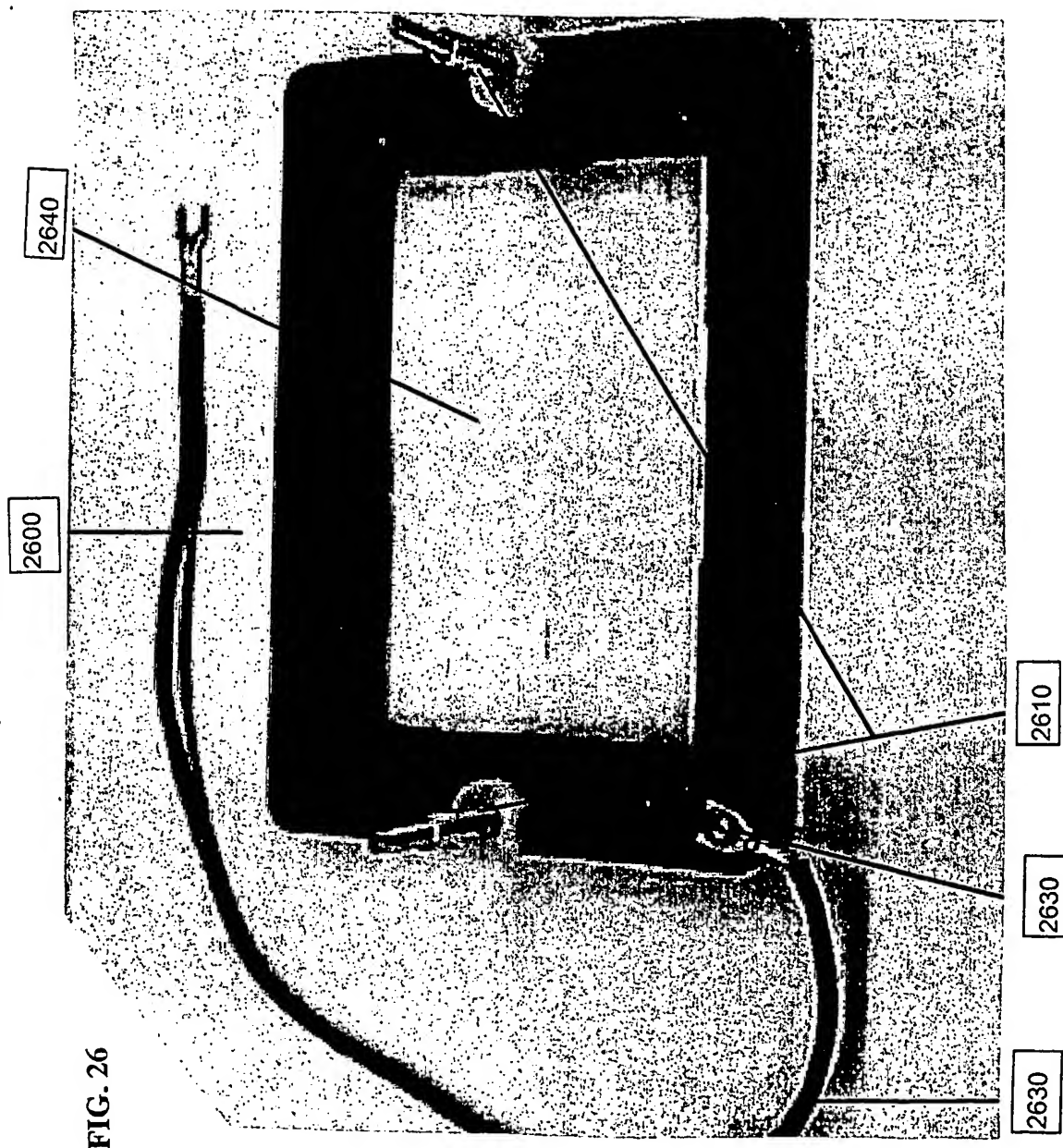


FIG. 27

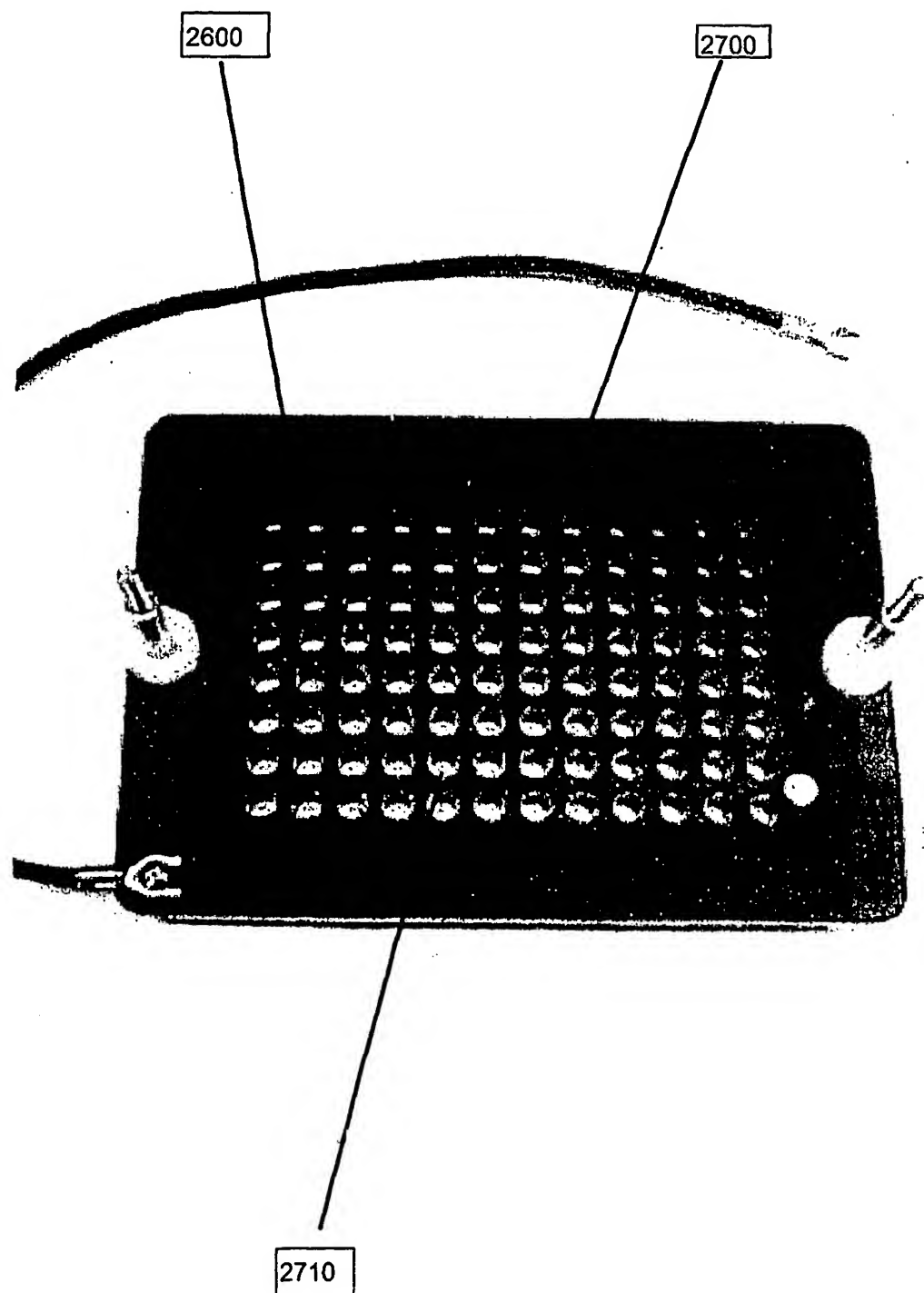
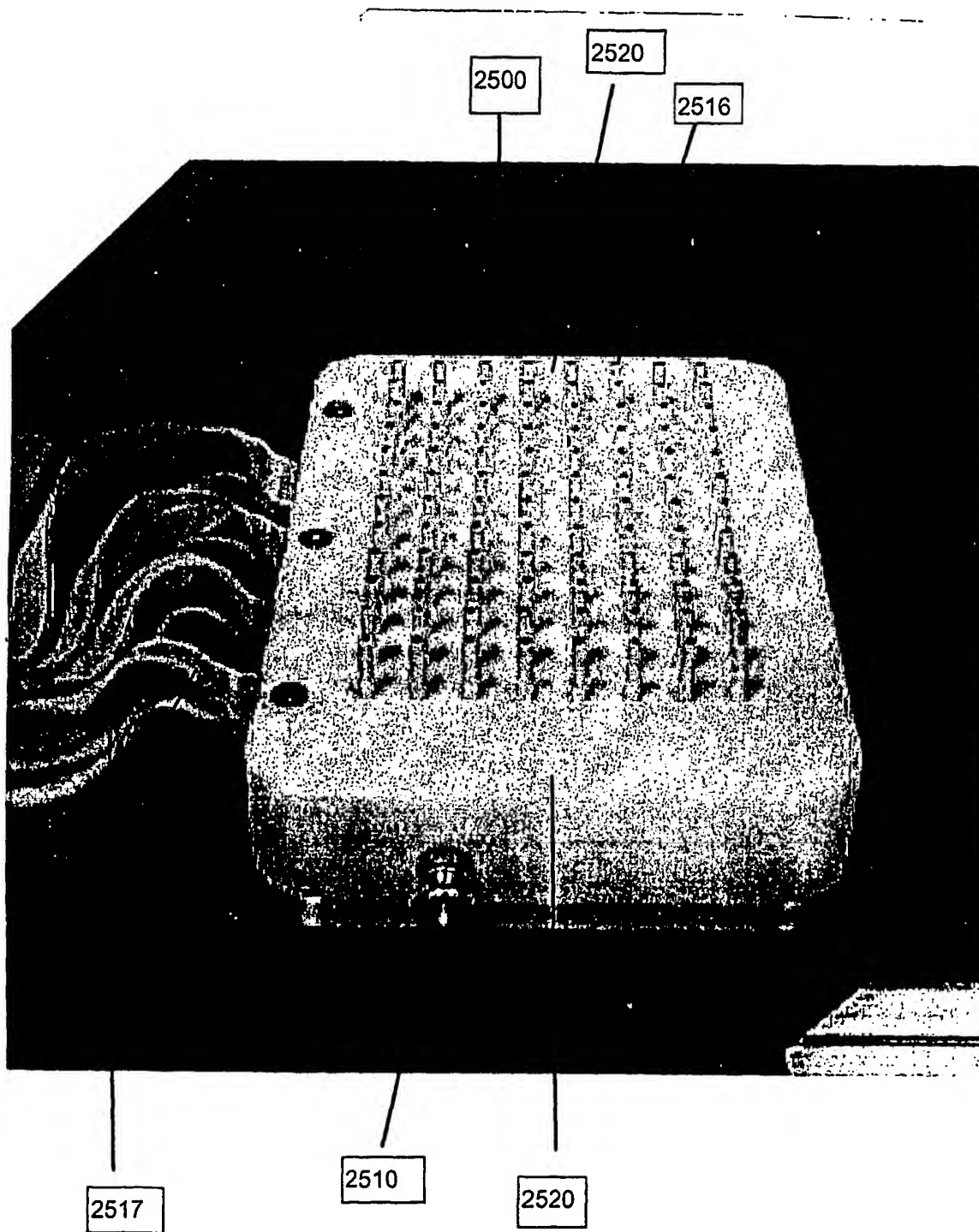


FIG. 25



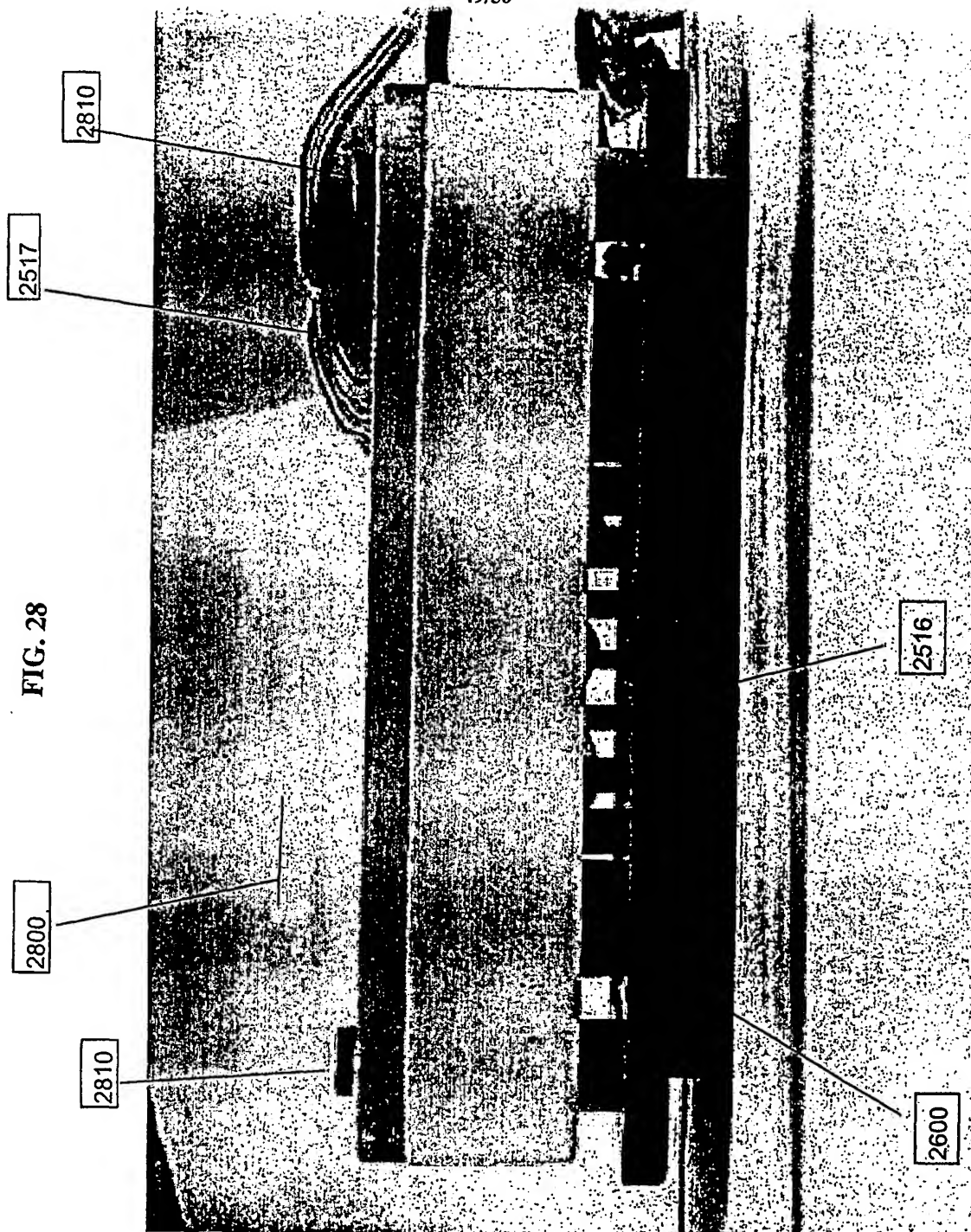


FIG. 29

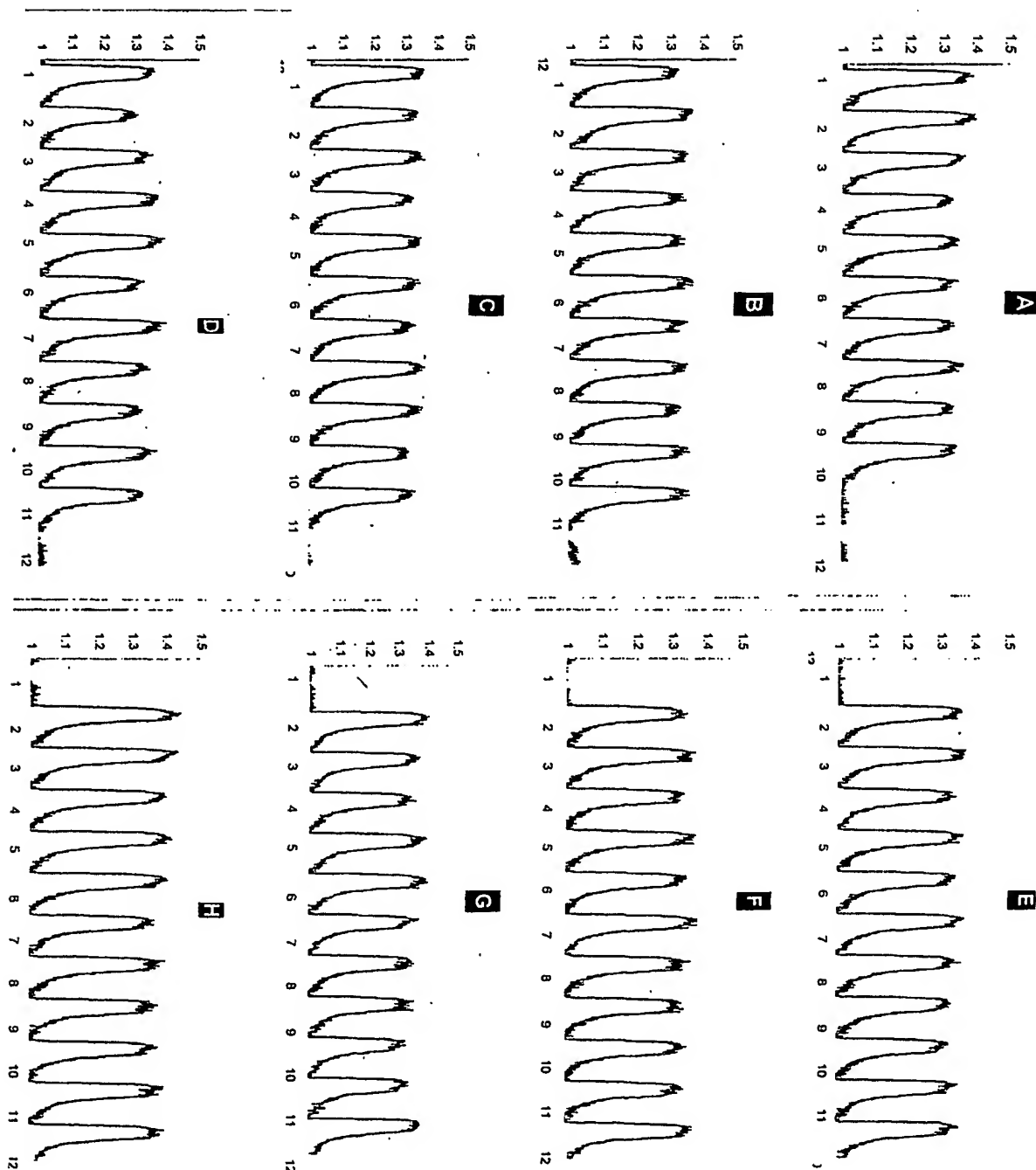


FIG. 30
Electric Field Stimulation of PN1 Cells Z=12
Testing 96-Well EFS Electrode System

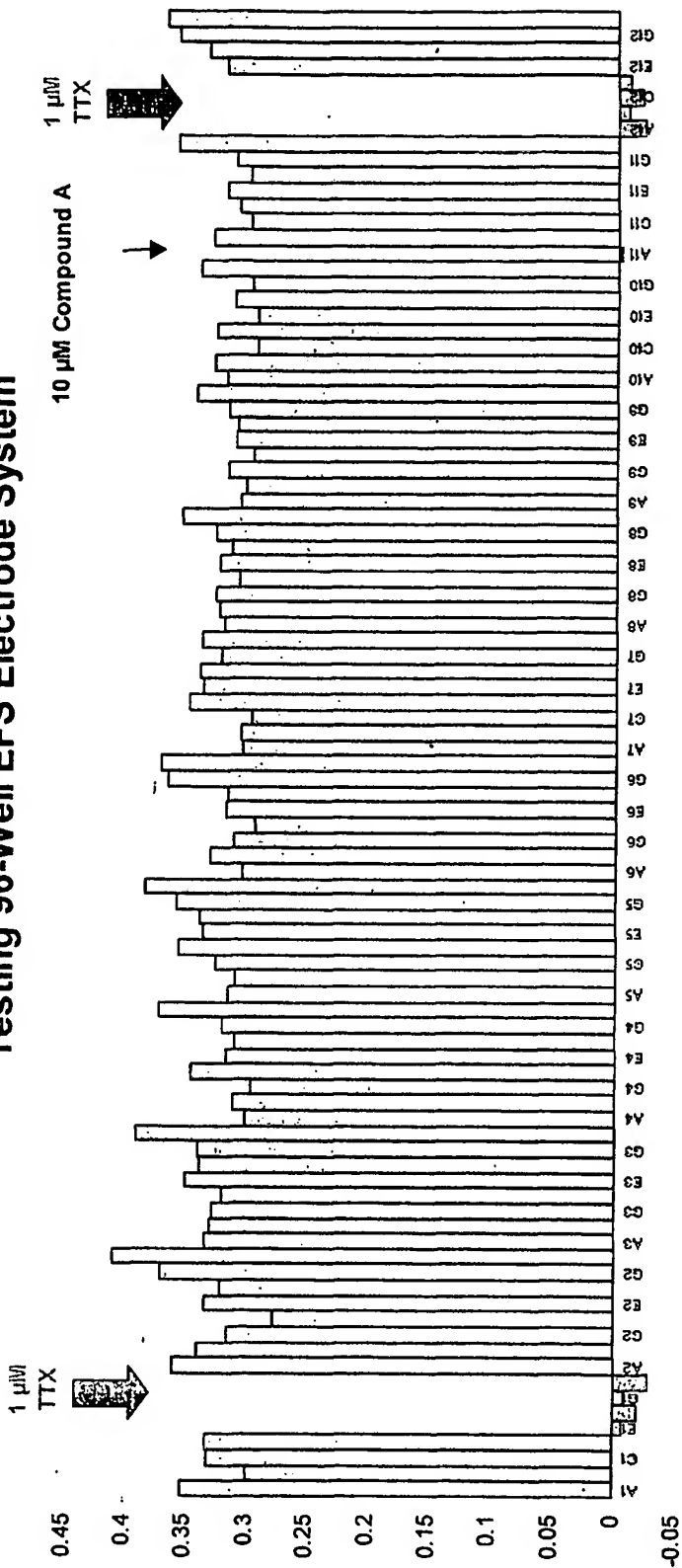


FIG. 31

Titration of Standards on PN1 cells stimulated with EFS

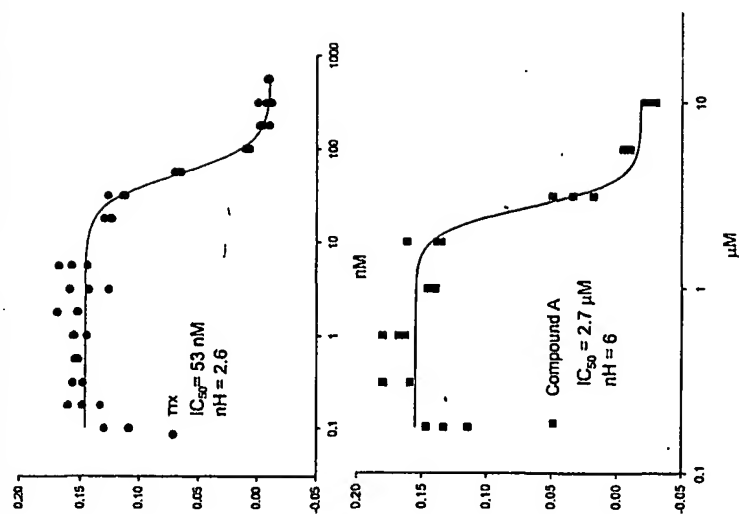


FIG. 32

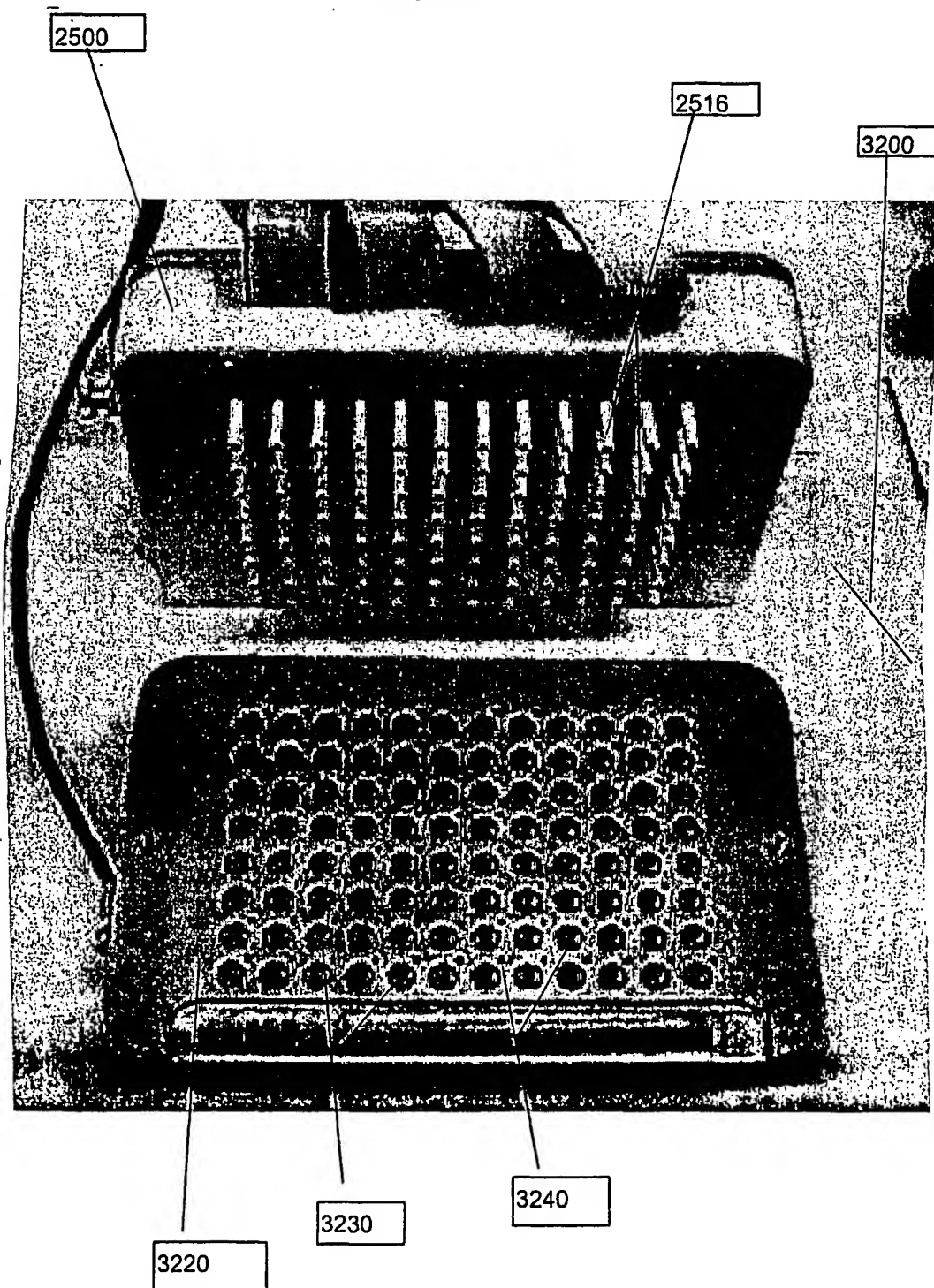


FIG. 33

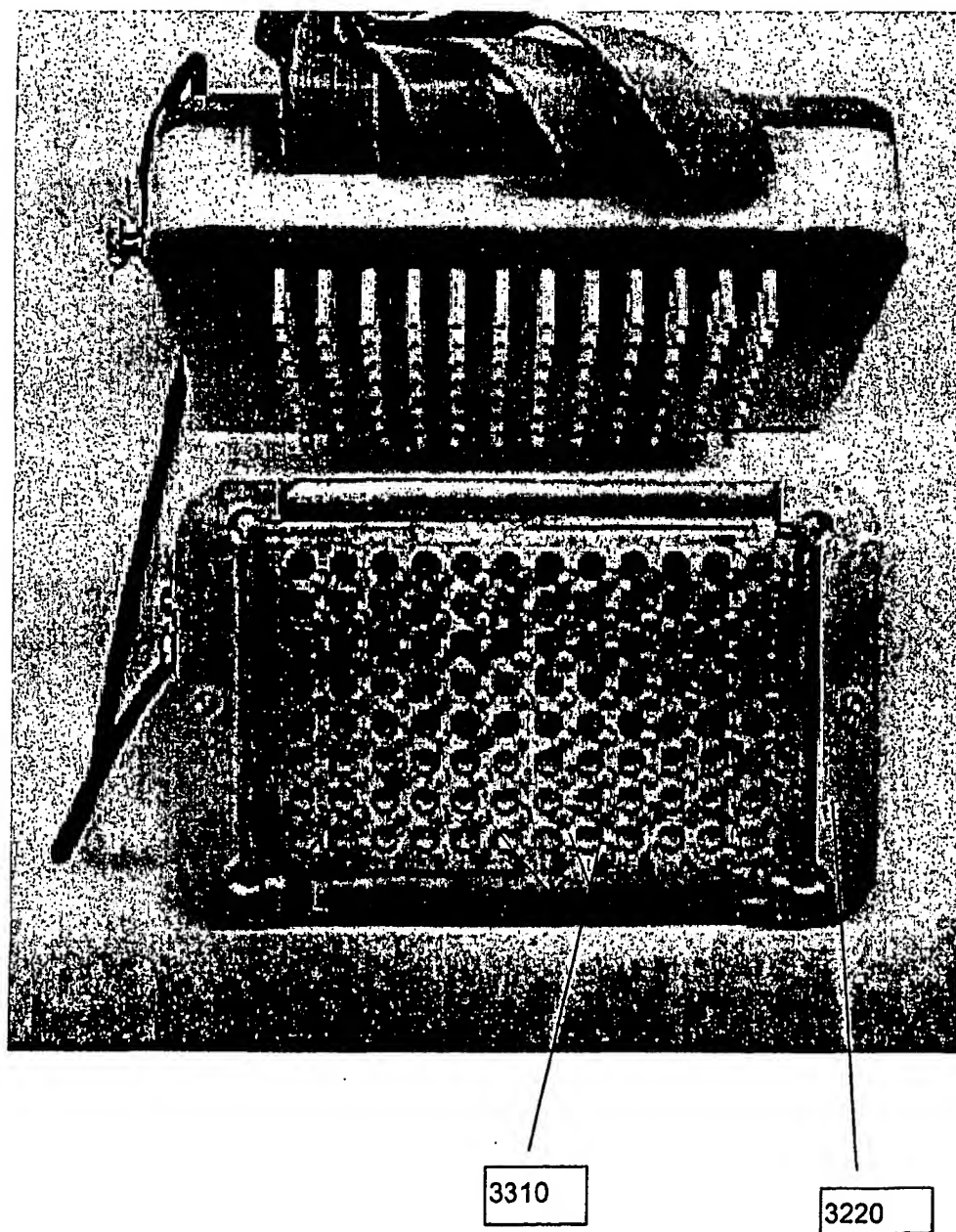


FIG. 34

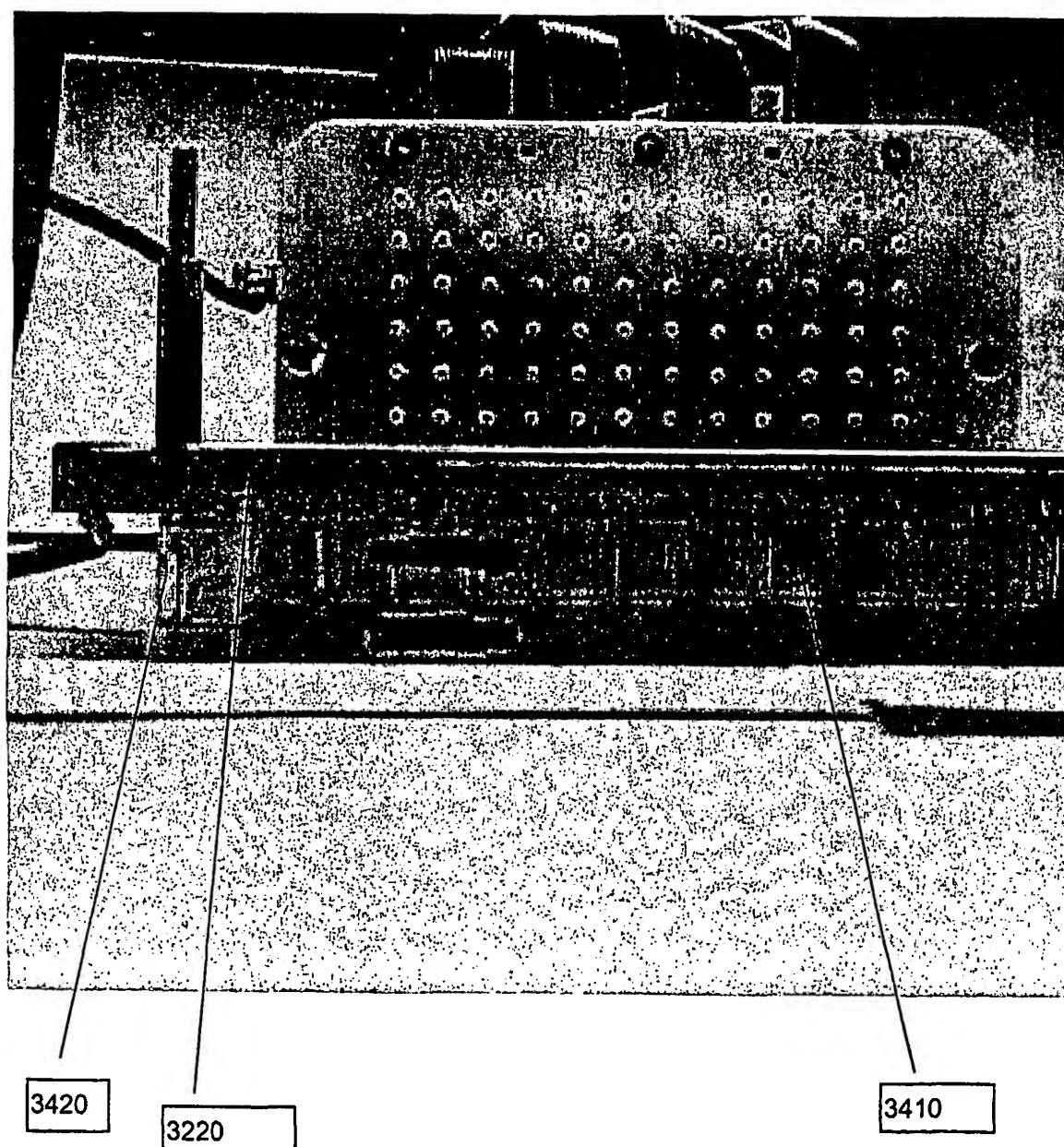
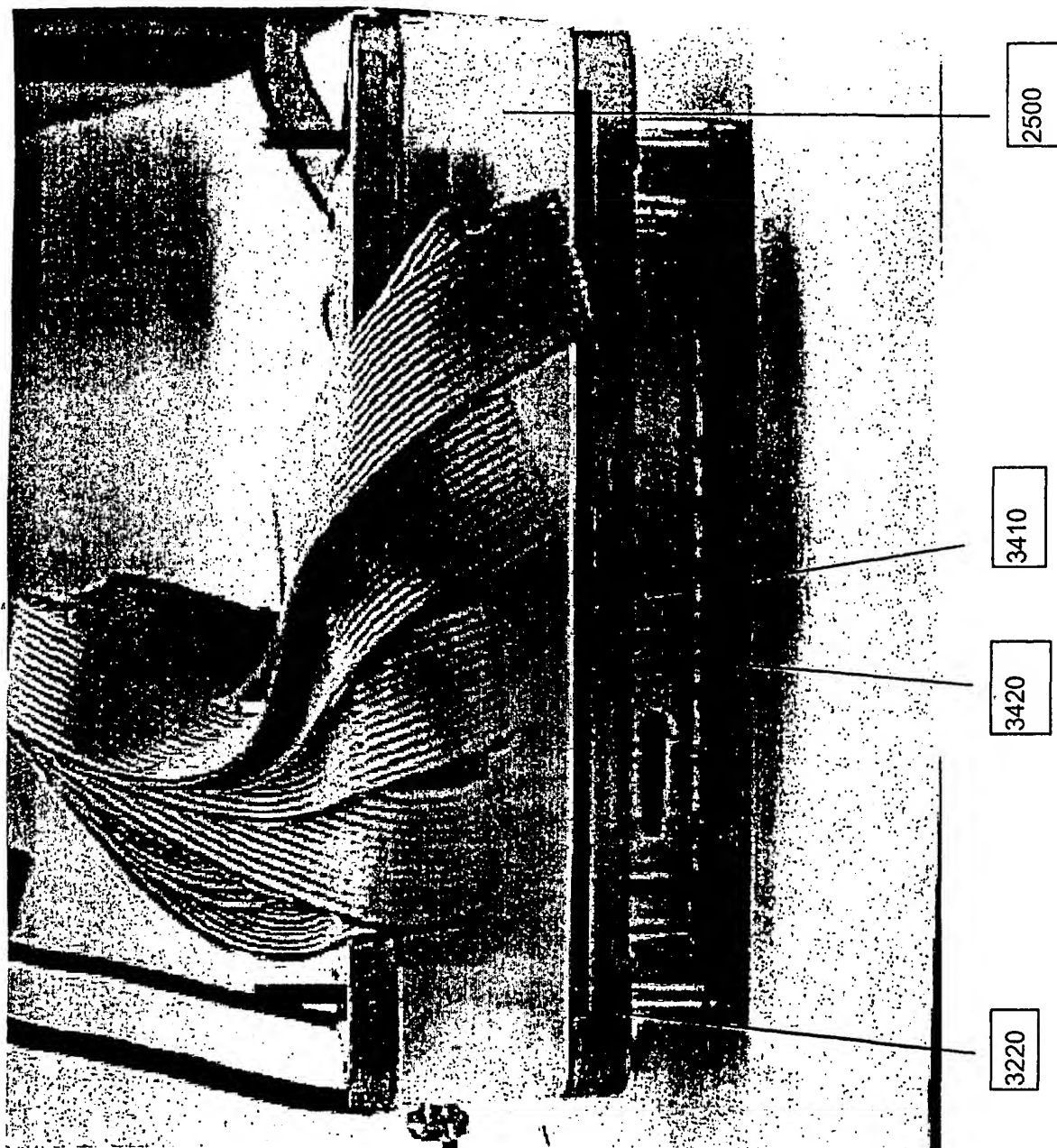


FIG. 35



SEQUENCE LISTING

<110> Kath, Gary S.
McManus, Owen
Garyantes, Tina
Bennett, Paul B., Jr.
Imredy, John P.
Augustine, Paul R.
Bugianesi, Randal M.

<120> ELECTRICAL FIELD STIMULATION OF
EUKARYOTIC CELLS

<130> 20794-PCT

<150> 60/304,955

<151> 2001-07-12

<160> 12

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 5874

<212> DNA

<213> Homo Sapiens

<400> 1

atggaattcc	ccattggatc	cctcgaaact	aacaacttcc	gtcgctttac	tccggagtca	60
ctgggtggaga	tagagaagca	aattgctgcc	aagcagggaa	caaagaaagc	cagagagaag	120
cataggggagc	agaaggacca	agaagagaag	cctcggcccc	agctggactt	gaaagcctgc	180
aaccagctgc	ccaagttcta	tggtgagctc	ccagcagaac	tgatcgggga	gccccctggag	240
gatctagatc	cgttctacag	cacacaccgg	acatttatgg	tgctgaacaa	agggaggacc	300
atttcccgggt	ttagtgccac	tcggggccctg	tggttattca	gtcctttcaa	cctgatcaga	360
agaacgggcca	tcaaagtgtc	tgtccactcg	tggttcagtt	tattttattac	ggtcactatt	420
ttgggttaatt	gtgtgtgcac	gaccogaact	gaccttccag	agaaaattga	atatgtcttc	480
actgtcattt	acacctttga	agccttgata	aagatactgg	caagaggatt	ttgtctaaat	540
gagttcacgt	acctgagaga	tccttggaa	tggttggtatt	ttagcgtcat	taccttgcca	600
tatgtttggca	cagcaataga	tctccgtggg	atctcaggcc	tgccggacatt	cagagttcct	660
agagcattaa	aaacagtttc	tgtgatccca	ggcctgaagg	tcattgtggg	ggccctgatt	720
cactcagtga	agaaactggc	tgatgtgacc	atcctcacca	tcttctgcct	aagtgttttt	780
gccttggtgg	ggctgcaact	cttcaagggc	aacctcaaaa	ataaatgtgt	caagaatgac	840
atggctgtca	atgagacaac	caactactca	tctcacagaa	aaccagatat	ctacataaat	900
aagcgaggca	cttctgaccc	cttactgtgt	ggcaatggat	ctgactcagg	ccactgcccc	960
gatggttata	tctgccttaa	aactttctgac	aaccgcggatt	ttaactacac	cagctttgat	1020
tccttttgctt	gggctttcct	ctcactgttc	cgcctcatga	cacaggattc	ctgggaacgc	1080
ctctaccagc	agaccctgag	gacttctggg	aaaatctata	tgatcttttt	tgtgctcgta	1140
atcttctctgg	gatctttcta	cctggtcaac	ttgatcttgg	ctgtagtcac	catggcgtat	1200
gaggagcaga	accaggcaac	cactgatgaa	attgaagcaa	aggagaagaa	gttccaggag	1260
gccctcgaga	tgctccggaa	ggagcaggag	gtgctagcag	cactagggat	tgacacaacc	1320
tctctccact	cccacaatgg	atcaccttta	acctccaaaa	atgccagtga	gagaaggcat	1380
agaataaaagc	caagagtgtc	agagggtccc	acagaagaca	acaaatcacc	ccgctctgat	1440
ccttacaacc	agcgaggat	gtcttttcta	ggcctgcct	ctggaaaacg	ccgggctagt	1500
catggcagtg	tgttccattt	ccgggtccct	ggccgagata	tctcactccc	tgagggagtc	1560
acagatgatg	gagtctttcc	tggagaccac	gaaagccatc	ggggctctct	gctgctgggt	1620
gggggtgctg	gccagcaagg	ccccctccct	agaagccctc	ttcctcaacc	cagcaaccct	1680
gactccaggc	atggagaaga	tgaacaccaa	ccgccgccca	ctagttagct	tgcccctgga	1740
gctgtcgatg	tctcggcatt	cgatgcagga	caaaagaaga	ctttcttgct	agcagaatac	1800
ttagatgaac	ctttccgggc	ccaaagggca	atgagtgttg	tcagtatcat	aacctccgtc	1860
cttgaggaac	tcgaggagtc	tgaacagaag	tgccccacct	gcttgaccag	cttgtctcag	1920

aagtatctga	tctgggattg	ctgccccatg	tgggtgaagc	tcaagacaat	tctctttggg	1980
cttgtgacgg	atcccttttg	agagctcacc	atcaccttgt	gcatcggtgg	gaacaccatc	2040
ttcatggcca	tggagcacca	tggcatgagc	cctaccttcg	aagccatgct	ccagataggc	2100
aacatcgtct	ttaccatatt	ttttactgct	gaaatgggtct	tcaaaatcat	tgccttcgac	2160
ccatactatt	atttccagaa	gaagtggaa	atctttgact	gcatcatcgt	cactgtgagt	2220
ctgctaggag	tgggctggc	caagaaggga	agcctgtctg	tgctgoggag	cttccgcttg	2280
ctggcgctat	tcaagctggc	caaatcctgg	cccaccttaa	acacactcat	caagatcatc	2340
ggaaactcag	tgggggcaact	ggggaacctc	accatcatcc	tggccatcat	tgtctttgtc	2400
tttgctctgg	ttggcaagca	gctcctaggg	gaaaactacc	gtaacaaccg	aaaaaatatc	2460
tccgcgcccc	atgaagactg	gccccgctgg	cacatgcacg	acttcttcca	ctctttcctc	2520
attgtcttcc	gtatcctctg	tggagagtgg	attgagaaca	tgtgggcctg	catggaagtt	2580
ggccaaaaat	ccatatgcct	catccttttc	ttgacggtga	tgggtgctagg	gaacctgggtg	2640
gtgcttaacc	tgttcatcgc	cctgctattg	aactctttca	gtgctgacaa	cctcacagcc	2700
ccggaggacg	tggtggagggt	gaacaacctg	cagggtggccc	tggcacggat	ccaggctctt	2760
ggccatcgta	ccaaacaggc	tctttgcagc	ttcttcagca	ggctcctgcc	attccccccag	2820
cccaaggcag	agcctgagct	ggtggtgaaa	ctcccactct	ccagctccaa	ggctgagaac	2880
cacattgctg	ccaacactgc	cagggggagc	tctggagggg	tccaagctcc	cagaggcccc	2940
agggatgag	acagtgaatt	catcgcta	ccgactgtgt	gggtctctgt	gcccatttgt	3000
gaggggtga	ctgactcttg	tgacttggag	gatgatgggt	gggaagatgc	tcagagcttc	3060
cagcaggaag	tgatccccaa	aggacagcag	gagcagctgc	agcaagtcca	gaggtgtggg	3120
gaccacctga	caccaggag	cccaggcact	ggaacatctt	ctgaggacct	ggctccatcc	3180
ctgggtgaga	cgtggaaaga	tgagtctgtt	cctcaggccc	ctgctgaggg	agtggacgac	3240
acaagctcct	ctgagggcag	cacggtggac	tgctatagtc	ctgaggaaat	cctgagggaag	3300
atccctgagc	tggcagatga	cctggaagaa	ccagatgact	gcttcacaga	aggatgcatt	3360
cgccactgtc	cctgctgcaa	actggatacc	accaagagtc	catgggatgt	gggctggcag	3420
gtgcgcgaag	cttgctaccg	tatcgtggag	cacagctggg	ttgagagctt	catcatcttc	3480
atgatcctgc	tcagcagtgg	atctctggcc	tttgaagact	attacctgga	ccagaagccc	3540
acgggtgaaag	ctttgctgga	gtacactgac	agggctcttca	cctttatctt	tgtgttcgag	3600
atgctgctta	agtgggtggc	ctatggcttc	aaaaagtact	tcaccaatgc	ctgggtgctgg	3660
ctggacttcc	tcatttgtgaa	tatctcactg	ataagctcca	cagcgaagat	tctggaatat	3720
tctgaagtgg	ctcccatcaa	agcccttcga	acccttcgcg	ctctgcgggc	actgagggct	3780
ctttctcgat	ttgaaggcat	gcggtgtgtg	gtggatgccc	tgggtgggcgc	catcccatcc	3840
atcatgaatg	tcctcctcgt	ctgcctcatc	ttctggctca	tcttcagcat	catgggtgtg	3900
aacctcttcg	cagggaagtt	ttggagggtg	atcaactata	ccgatggaga	gttttccctt	3960
gtacctttgt	cgatttgtgaa	taacaagtct	gactgcaaga	ttcaaaactc	cactggcagc	4020
ttcttctggg	tcaatgtgaa	agtcaacttt	gataatgttg	caatgggtta	ccttgcactt	4080
ctgcagggtg	caacctttta	aggctggatg	gacattatgt	atgcagctgt	tgattcccgg	4140
gaggtcaaca	tgcaacccaa	gtgggaggac	aacgtgtaca	tgtattttgt	ctttgtcatc	4200
ttcatcattt	ttggaggctt	cttcacactg	aatctctttg	ttgggtcat	aattgacaac	4260
ttcaatcaac	agaaaaaaa	gttagggggc	caggacatct	tcatgacaga	ggagcagaag	4320
aaatactaca	atgccatgaa	gaagtgtggc	tccaagaagc	cccagaagcc	catcccacgg	4380
cccctggaca	agttccaggg	ttttgtcttt	gacatctgta	ccagacaagc	ttttgacatc	4440
accatcatgg	tcctcatctg	cctcaacatg	atcaccatga	tgggtggagac	tgatgaccac	4500
agtgaagaaa	agacgaaaat	tctgggcaaa	atcaaccagt	tctttgtggc	cgtcttcaca	4560
ggcgaatgtg	tcatgaagat	gttcgctttg	aggcagtact	acttcacaaa	tggctggaat	4620
gtgtttgact	tcatttgtgt	ggttctctcc	attgcgagcc	tgattttttc	tgcaattctt	4680
aagtcacttc	aaagttactt	ctccccaaag	ctcttcagag	tcacccgcct	ggccccgaat	4740
ggccgcatcc	tcagactgat	ccgagcggcc	aaggggatcc	gcacactgct	ctttgccttc	4800
atgatgtccc	tgcctgccct	cttcaacatc	gggctgttgc	tattccttgt	catgttcac	4860
tactccatct	tcggtatgtc	cagctttccc	catgtgaggt	gggaggctgg	catcgacgac	4920
atgttcaact	tccagacctt	cgccaacagc	atgctgtgcc	tcttccagat	taccacgtcg	4980
gccggctggg	atggcctcct	cagccccatc	ctcaacacag	ggccccctta	ctgtgacccc	5040
aatctgcccc	acagcaatgg	caccagaggg	gactgtggga	gcccagccgt	aggcatcatc	5100
ttcttcacca	cctacatcat	catctccttc	ctcatcgtgg	tcaacatgta	cattgcagtg	5160
attctggaga	atttcaatgt	ggccacggag	gagagcactg	agcctctgag	tgaggacgac	5220
tttgacatgt	tctatgagac	ctgggagaag	tttgacccag	aggccactca	gtttattacc	5280
ttttctgctc	tctcggactt	tgcagacact	ctctctggtc	ccctgagaat	cccaaaaccc	5340
aatcgaataa	tactgatcca	gatggacctg	cctttgtgtc	ctggagataa	gatccactgc	5400
ttggacatcc	tttttgcttt	caccaagaat	gtcctaggag	aatccgggga	gttggattct	5460
ctgaaggcaa	atatggagga	gaagtttatg	gcaactaatc	tttcaaaatc	atcctatgaa	5520
ccaatagcaa	ccactctccg	atggaagcaa	gaagacattt	cagccactgt	cattcaaaag	5580
gcctatcgga	gctatgtgct	gcaccgctcc	atggcactct	ctaacacccc	atgtgtgccc	5640


```

agagctgagg aggaggctgc atcactccca gatgaaggtt ttgttgcatc cacagcaaat 5700
gaaaattgtg tactcccaga caaatctgaa actgcttctg ccacatcatt cccaccgtcc 5760
tatgagagtg tcactagagg ccttagtgat agagtcaaca tgaggacatc tagctcaata 5820
caaaatgaag atgaagccac cagtatggag ctgattgccc ctggggcccta gtga 5874

```

<210> 2
 <211> 1956
 <212> PRT
 <213> Homo Sapiens

<400> 2

Met	Glu	Phe	Pro	Ile	Gly	Ser	Leu	Glu	Thr	Asn	Asn	Phe	Arg	Arg	Phe	1	5	10	15
Thr	Pro	Glu	Ser	Leu	Val	Glu	Ile	Glu	Lys	Gln	Ile	Ala	Ala	Lys	Gln	20	25	30	35
Gly	Thr	Lys	Lys	Ala	Arg	Glu	Lys	His	Arg	Glu	Gln	Lys	Asp	Gln	Glu	40	45	50	55
Glu	Lys	Pro	Arg	Pro	Gln	Leu	Asp	Leu	Lys	Ala	Cys	Asn	Gln	Leu	Pro	60	65	70	75
Lys	Phe	Tyr	Gly	Glu	Leu	Pro	Ala	Glu	Leu	Ile	Gly	Glu	Pro	Leu	Glu	80	85	90	95
Asp	Leu	Asp	Pro	Phe	Tyr	Ser	Thr	His	Arg	Thr	Phe	Met	Val	Leu	Asn	100	105	110	115
Lys	Gly	Arg	Thr	Ile	Ser	Arg	Phe	Ser	Ala	Thr	Arg	Ala	Leu	Trp	Leu	120	125	130	135
Phe	Ser	Pro	Phe	Asn	Leu	Ile	Arg	Arg	Thr	Ala	Ile	Lys	Val	Ser	Val	140	145	150	155
His	Ser	Trp	Phe	Ser	Leu	Phe	Ile	Thr	Val	Thr	Ile	Leu	Val	Asn	Cys	160	165	170	175
Val	Cys	Met	Thr	Arg	Thr	Asp	Leu	Pro	Glu	Lys	Ile	Glu	Tyr	Val	Phe	180	185	190	195
Thr	Val	Ile	Tyr	Thr	Phe	Glu	Ala	Leu	Ile	Lys	Ile	Leu	Ala	Arg	Gly	200	205	210	215
Phe	Cys	Leu	Asn	Glu	Phe	Thr	Tyr	Leu	Arg	Asp	Pro	Trp	Asn	Trp	Leu	220	225	230	235
Asp	Phe	Ser	Val	Ile	Thr	Leu	Ala	Tyr	Val	Gly	Thr	Ala	Ile	Asp	Leu	240	245	250	255
Arg	Gly	Ile	Ser	Gly	Leu	Arg	Thr	Phe	Arg	Val	Leu	Arg	Ala	Leu	Lys	260	265	270	275
Thr	Val	Ser	Val	Ile	Pro	Gly	Leu	Lys	Val	Ile	Val	Gly	Ala	Leu	Ile	280	285	290	295
His	Ser	Val	Lys	Lys	Leu	Ala	Asp	Val	Thr	Ile	Leu	Thr	Ile	Phe	Cys	300	305	310	315
Leu	Ser	Val	Phe	Ala	Leu	Val	Gly	Leu	Gln	Leu	Phe	Lys	Gly	Asn	Leu	320	325	330	335
Lys	Asn	Lys	Cys	Val	Lys	Asn	Asp	Met	Ala	Val	Asn	Glu	Thr	Thr	Asn	340	345	350	355
Tyr	Ser	Ser	His	Arg	Lys	Pro	Asp	Ile	Tyr	Ile	Asn	Lys	Arg	Gly	Thr	360	365	370	375
Ser	Asp	Pro	Leu	Leu	Cys	Gly	Asn	Gly	Ser	Asp	Ser	Gly	His	Cys	Pro	380	385	390	395
Asp	Gly	Tyr	Ile	Cys	Leu	Lys	Thr	Ser	Asp	Asn	Pro	Asp	Phe	Asn	Tyr	400	405	410	415
Thr	Ser	Phe	Asp	Ser	Phe	Ala	Trp	Ala	Phe	Leu	Ser	Leu	Phe	Arg	Leu	420	425	430	435
Met	Thr	Gln	Asp	Ser	Trp	Glu	Arg	Leu	Tyr	Gln	Gln	Thr	Leu	Arg	Thr	440	445	450	455
Ser	Gly	Lys	Ile	Tyr	Met	Ile	Phe	Phe	Val	Leu	Val	Ile	Phe	Leu	Gly	460	465	470	475
Ser	Phe	Tyr	Leu	Val	Asn	Leu	Ile	Leu	Ala	Val	Val	Thr	Met	Ala	Tyr	480	485	490	495

385					390					395				400
Glu	Glu	Gln	Asn	Gln	Ala	Thr	Thr	Asp	Glu	Ile	Glu	Ala	Lys	Glu
				405					410					415
Lys	Phe	Gln	Glu	Ala	Leu	Glu	Met	Leu	Arg	Lys	Glu	Gln	Glu	Val
			420					425						430
Ala	Ala	Leu	Gly	Ile	Asp	Thr	Thr	Leu	His	Ser	His	Asn	Gly	Ser
		435					440				445			
Pro	Leu	Thr	Ser	Lys	Asn	Ala	Ser	Glu	Arg	Arg	His	Arg	Ile	Lys
	450					455					460			Pro
Arg	Val	Ser	Glu	Gly	Ser	Thr	Glu	Asp	Asn	Lys	Ser	Pro	Arg	Ser
	465				470					475				480
Pro	Tyr	Asn	Gln	Arg	Arg	Met	Ser	Phe	Leu	Gly	Leu	Ala	Ser	Gly
				485				490						495
Arg	Arg	Ala	Ser	His	Gly	Ser	Val	Phe	His	Phe	Arg	Ser	Pro	Gly
			500					505					510	Arg
Asp	Ile	Ser	Leu	Pro	Glu	Gly	Val	Thr	Asp	Asp	Gly	Val	Phe	Pro
	515						520					525		Gly
Asp	His	Glu	Ser	His	Arg	Gly	Ser	Leu	Leu	Leu	Gly	Gly	Gly	Ala
	530					535					540			Gly
Gln	Gln	Gly	Pro	Leu	Pro	Arg	Ser	Pro	Leu	Pro	Gln	Pro	Ser	Asn
	545				550					555				560
Asp	Ser	Arg	His	Gly	Glu	Asp	Glu	His	Gln	Pro	Pro	Pro	Thr	Ser
				565				570						575
Leu	Ala	Pro	Gly	Ala	Val	Asp	Val	Ser	Ala	Phe	Asp	Ala	Gly	Gln
			580				585						590	Lys
Lys	Thr	Phe	Leu	Ser	Ala	Glu	Tyr	Leu	Asp	Glu	Pro	Phe	Arg	Ala
	595						600					605		Gln
Arg	Ala	Met	Ser	Val	Val	Ser	Ile	Ile	Thr	Ser	Val	Leu	Glu	Glu
	610					615					620			Leu
Glu	Glu	Ser	Glu	Gln	Lys	Cys	Pro	Pro	Cys	Leu	Thr	Ser	Leu	Ser
	625				630					635				Gln
Lys	Tyr	Leu	Ile	Trp	Asp	Cys	Cys	Pro	Met	Trp	Val	Lys	Leu	Lys
				645					650					655
Ile	Leu	Phe	Gly	Leu	Val	Thr	Asp	Pro	Phe	Ala	Glu	Leu	Thr	Ile
			660					665					670	Thr
Leu	Cys	Ile	Val	Val	Asn	Thr	Ile	Phe	Met	Ala	Met	Glu	His	His
	675					680					685			Gly
Met	Ser	Pro	Thr	Phe	Glu	Ala	Met	Leu	Gln	Ile	Gly	Asn	Ile	Val
	690					695					700			Phe
Thr	Ile	Phe	Phe	Thr	Ala	Glu	Met	Val	Phe	Lys	Ile	Ile	Ala	Phe
	705				710					715				Asp
Pro	Tyr	Tyr	Tyr	Phe	Gln	Lys	Lys	Trp	Asn	Ile	Phe	Asp	Cys	Ile
				725					730					735
Val	Thr	Val	Ser	Leu	Leu	Glu	Leu	Gly	Val	Ala	Lys	Lys	Gly	Ser
			740					745					750	Leu
Ser	Val	Leu	Arg	Ser	Phe	Arg	Leu	Leu	Arg	Val	Phe	Lys	Leu	Ala
	755						760				765			Lys
Ser	Trp	Pro	Thr	Leu	Asn	Thr	Leu	Ile	Lys	Ile	Ile	Gly	Asn	Ser
	770					775					780			Val
Gly	Ala	Leu	Gly	Asn	Leu	Thr	Ile	Ile	Leu	Ala	Ile	Ile	Val	Phe
	785				790					795				Val
Phe	Ala	Leu	Val	Gly	Lys	Gln	Leu	Leu	Gly	Glu	Asn	Tyr	Arg	Asn
				805					810					815
Arg	Lys	Asn	Ile	Ser	Ala	Pro	His	Glu	Asp	Trp	Pro	Arg	Trp	His
				820				825					830	Met
His	Asp	Phe	Phe	His	Ser	Phe	Leu	Ile	Val	Phe	Arg	Ile	Leu	Cys
	835					840					845			Gly
Glu	Trp	Ile	Glu	Asn	Met	Trp	Ala	Cys	Met	Glu	Val	Gly	Gln	Lys
	850					855					860			Ser
Ile	Cys	Leu	Ile	Leu	Phe	Leu	Thr	Val	Met	Val	Leu	Gly	Asn	Leu
	865				870				875					880
Val	Leu	Asn	Leu	Phe	Ile	Ala	Leu	Leu	Leu	Asn	Ser	Phe	Ser	Ala

				885					890					895	
Asn	Leu	Thr	Ala	Pro	Glu	Asp	Asp	Gly	Glu	Val	Asn	Asn	Leu	Gln	Val
			900					905					910		
Ala	Leu	Ala	Arg	Ile	Gln	Val	Phe	Gly	His	Arg	Thr	Lys	Gln	Ala	Leu
		915					920					925			
Cys	Ser	Phe	Phe	Ser	Arg	Ser	Cys	Pro	Phe	Pro	Gln	Pro	Lys	Ala	Glu
	930					935					940				
Pro	Glu	Leu	Val	Val	Lys	Leu	Pro	Leu	Ser	Ser	Ser	Lys	Ala	Glu	Asn
945					950					955				960	
His	Ile	Ala	Ala	Asn	Thr	Ala	Arg	Gly	Ser	Ser	Gly	Gly	Leu	Gln	Ala
			965					970						975	
Pro	Arg	Gly	Pro	Arg	Asp	Glu	His	Ser	Asp	Phe	Ile	Ala	Asn	Pro	Thr
			980					985					990		
Val	Trp	Val	Ser	Val	Pro	Ile	Ala	Glu	Gly	Glu	Ser	Asp	Leu	Asp	Asp
	995						1000					1005			
Leu	Glu	Asp	Asp	Gly	Gly	Glu	Asp	Ala	Gln	Ser	Phe	Gln	Gln	Glu	Val
	1010					1015						1020			
Ile	Pro	Lys	Gly	Gln	Gln	Glu	Gln	Leu	Gln	Gln	Val	Glu	Arg	Cys	Gly
1025					1030					1035					1040
Asp	His	Leu	Thr	Pro	Arg	Ser	Pro	Gly	Thr	Gly	Thr	Ser	Ser	Glu	Asp
				1045					1050					1055	
Leu	Ala	Pro	Ser	Leu	Gly	Glu	Thr	Trp	Lys	Asp	Glu	Ser	Val	Pro	Gln
		1060						1065					1070		
Ala	Pro	Ala	Glu	Gly	Val	Asp	Asp	Thr	Ser	Ser	Ser	Glu	Gly	Ser	Thr
	1075						1080					1085			
Val	Asp	Cys	Leu	Asp	Pro	Glu	Ile	Leu	Arg	Lys	Ile	Pro	Glu	Leu	
	1090					1095					1100				
Ala	Asp	Asp	Leu	Glu	Glu	Pro	Asp	Asp	Cys	Phe	Thr	Glu	Gly	Cys	Ile
1105					1110					1115					1120
Arg	His	Cys	Pro	Cys	Cys	Lys	Leu	Asp	Thr	Thr	Lys	Ser	Pro	Trp	Asp
			1125						1130					1135	
Val	Gly	Trp	Gln	Val	Arg	Lys	Thr	Cys	Tyr	Arg	Ile	Val	Glu	His	Ser
		1140						1145					1150		
Trp	Phe	Glu	Ser	Phe	Ile	Ile	Phe	Met	Ile	Leu	Leu	Ser	Ser	Gly	Ser
	1155						1160					1165			
Leu	Ala	Phe	Glu	Asp	Tyr	Tyr	Leu	Asp	Gln	Lys	Pro	Thr	Val	Lys	Ala
	1170				1175						1180				
Leu	Leu	Glu	Tyr	Thr	Asp	Arg	Val	Phe	Thr	Phe	Ile	Phe	Val	Phe	Glu
1185					1190					1195					1200
Met	Leu	Leu	Lys	Trp	Val	Ala	Tyr	Gly	Phe	Lys	Lys	Tyr	Phe	Thr	Asn
			1205						1210					1215	
Ala	Trp	Cys	Trp	Leu	Asp	Phe	Leu	Ile	Val	Asn	Ile	Ser	Leu	Ile	Ser
		1220						1225					1230		
Leu	Thr	Ala	Lys	Ile	Leu	Glu	Tyr	Ser	Glu	Val	Ala	Pro	Ile	Lys	Ala
	1235						1240					1245			
Leu	Arg	Thr	Leu	Arg	Ala	Leu	Arg	Pro	Leu	Arg	Ala	Leu	Ser	Arg	Phe
	1250				1255						1260				
Glu	Gly	Met	Arg	Val	Val	Val	Asp	Ala	Leu	Val	Gly	Ala	Ile	Pro	Ser
1265					1270					1275					1280
Ile	Met	Asn	Val	Leu	Leu	Val	Cys	Leu	Ile	Phe	Trp	Leu	Ile	Phe	Ser
			1285						1290					1295	
Ile	Met	Gly	Val	Asn	Leu	Phe	Ala	Gly	Lys	Phe	Trp	Arg	Cys	Ile	Asn
		1300						1305					1310		
Tyr	Thr	Asp	Gly	Glu	Phe	Ser	Leu	Val	Pro	Leu	Ser	Ile	Val	Asn	Asn
	1315						1320					1325			
Lys	Ser	Asp	Cys	Lys	Ile	Gln	Asn	Ser	Thr	Gly	Ser	Phe	Phe	Trp	Val
	1330				1335						1340				
Asn	Val	Lys	Val	Asn	Phe	Asp	Asn	Val	Ala	Met	Gly	Tyr	Leu	Ala	Leu
1345					1350					1355					1360
Leu	Gln	Val	Ala	Thr	Phe	Lys	Gly	Trp	Met	Asp	Ile	Met	Tyr	Ala	Ala
			1365						1370					1375	
Val	Asp	Ser	Arg	Glu	Val	Asn	Met	Gln	Pro	Lys	Trp	Glu	Asp	Asn	Val

- 6 -

1875 1880 1885
 Leu Pro Asp Glu Gly Phe Val Ala Phe Thr Ala Asn Glu Asn Cys Val
 1890 1895 1900
 Leu Pro Asp Lys Ser Glu Thr Ala Ser Ala Thr Ser Phe Pro Pro Ser
 1905 1910 1915 1920
 Tyr Glu Ser Val Thr Arg Gly Leu Ser Asp Arg Val Asn Met Arg Thr
 1925 1930 1935
 Ser Ser Ser Ile Gln Asn Glu Asp Glu Ala Thr Ser Met Glu Leu Ile
 1940 1945 1950
 Ala Pro Gly Pro
 1955

<210> 3
 <211> 7898
 <212> DNA
 <213> Homo Sapiens

<400> 3
 cgaggccgcc gccgtgcgct ccgcccgggcy agccggagcc ggagtcgagc cgcggccggg 60
 agccgggcyg gctgggggacg cgggcccggg gggaggcgcc tggggggccg ggccggggcc 120
 gggggccgag gcgctggggg cgggggcccg ggcggggcgc cgagcggggt ccgcggtgac 180
 cgcgccgccc gggcgatgcc cgcggggacg ccgcccggca gcagagcgag gtgctgccgg 240
 ccgccaccat gaccgagggc gcacggggcg ccgacgaggt ccgggtgccc ctgggcccgc 300
 cgcctctggt ccctgcggcg ttggtggggg cgtccccgga gagccccggg gcgcccggac 360
 gcgagggcga gcggggggtcc gagctcgggc tgtcacctcc cgagagcccc gcgggccgagc 420
 gcgggcgcgga gctgggtgcc gacgaggagc agcgcgctcc gtaccgggcc ttggcgggca 480
 cggctctctt ctgcctcggt cagaccacgc ggcgcgcgag ctgggtgctc cggctgggtc 540
 gcaacccatg gttcgagcac gtgagcatgc tggtaatcat gctcaactgc gtgaccctgg 600
 gcatgttcg gcccgtgtgag gacgttgagt gcggtccgca gcgctgcaac atcctggagg 660
 cctttgacgc cttcattttc gccttttttg cgggtggagt ggtcatcaag atggtggcct 720
 tggggctggt cgggcagaag tgttacctgg gtgacacgtg gaacaggctg gatttcttca 780
 tcgtcgtggc gggcatgatg gactactcgt tggacggaca caacgtgagc ctctcggtca 840
 tcaggaccgt gcggtgtgct cggccccctc gcgccatcaa ccgctgctc agcatgcgga 900
 tcttggctac tctgctgctg gatacgtgct ccatgctcgg gaacgtcctt ctgctgtgct 960
 tcttgcctct cttcattttc ggcatcggtt ggcacacgtc ctgggctggc ctctcttcca 1020
 accgctgctt cctggacagt gcctttgtca ggaacaacaa cctgaccttc ctgcccgcgt 1080
 actaccagac ggaggagggt gaggagaacc cgttcatctg ctctcacgc cgagacaacg 1140
 gcatgcagaa gtgctcgcac atccccggcc gccgcgagct ggcgatgccc tgcacctgg 1200
 gctgggaggg ctacacgcag ccgcaggcag agggggtggg cgctgcacgc aacgcttca 1260
 tcaactggaa ccagtactac aacgtgtgcc gctcgggtga ctccaacccc cacaacggtg 1320
 ccatcaactt cgacaacatc ggctacgctt ggattgccat ctccaggtg atcacgctgg 1380
 aaggctgggt ggacatcatg tactacgtca tggacgcccc ctcttctac aacttcatct 1440
 atttcatcct gctcatcatc gtgggtcct tcttcatgat caacctgtgc ctggtggtga 1500
 ttgccacgca gttctcgag acgaagcagc gggagagtca gctgatgcgg gagcagcggg 1560
 cagccaccct gtccaacgac agcacgctgg ccagcttctc cgagcctggc agctgctacg 1620
 aagagctgct gaagtacgtg ggccacatat tccgcaaggt caagcggcgc agcttgcgcc 1680
 tctacgcccg ctggcagagc cgctggcgca agaagtgga cccagtgct gtgcaaggcc 1740
 aggggtcccg gcaccgccag cgcggggcag gcaggcacac agcctcgggt caccacctgg 1800
 tctaccacca ccatcaccac caccaccacc actaccattt cagccatggc agcccccgca 1860
 ggccccggcc cgagccaggc gcctgcgaca ccaggctggt ccgagctggc ggcccccct 1920
 cgccaccttc ccaggccgc ggacccccg acgcagagtc tgtgcacagc atctaccatg 1980
 ccgactgcca catagagggg ccgcaggaga gggccccggg ggcacatgcc gcagccactg 2040
 ccgctgccag cctcaggctg gccacagggt tgggacccat gaactacccc acgatcctgc 2100
 cctcagggtt gggcagcggc aaaggcagca ccaggccccg acccaagggg aagtgggccg 2160
 gtggaccgcc aggcaccggg gggcacggcc cgttgagctt gaacagccct gatccctacg 2220
 agaagatccc ccatgtggtc ggggagcatg gactgggcca ggccccctgg catctgtcgg 2280
 gcctcagtg gtccctgccc ctgcccagcc cccagcggg cacactgacc tgtgagctga 2340
 agagctgccc gtactgcacc cgtgccttgg aggacccgga ggggtgagtc agcggctcgg 2400
 aaagtggaga ctcagatggc cgtggcgtct atgaattcac gcaggacgtc cggcacgggtg 2460
 accgctggga ccccacgcga ccaccccggt cgacggacac accaggccca ggcccaggca 2520
 gccccagcg gcgggcacag cagagggcag ccccgggcga gccaggctgg atgggcccgc 2580
 tctgggttac cttcagcggc aagctgcgcc gcatcgtgga cagcaagtag ttcagccgtg 2640

gcatcatgat	ggccatcctt	gtcaaacacgc	tgagcatggg	cgtggagtag	catgagcagc	2700
ccgaggagct	gactaatgct	ctggagatca	gcaacatcgt	gttcaccagc	atgtttgccc	2760
tggagatgct	gctgaagctg	ctggcctgcg	gccctctggg	ctacatccgg	aaccctgaca	2820
acatcttcga	cggcatcatc	gtggtcatca	gcgtctggga	gatcgtgggg	caggcggacg	2880
gtggcttgct	tgtgctgcgc	accttccggc	tgctgcgtgt	gctgaagctg	gtgcgctttc	2940
tgccagccct	gcggcgccag	ctcgtggtgc	tggatgaagac	catggacaac	gtggctacct	3000
tctgcacgct	gctcatgctc	ttcattttca	tcttcagcat	cctgggcatg	caccttttctg	3060
gctgcaagtt	cagcctgaag	acagacaccg	gagacaccgt	gcctgacagg	aagaacttcg	3120
actccctgct	gtggggccatc	gtcaccgtgt	tccagatcct	gaccacaggag	gactggaacg	3180
tggctcctgta	caacggcatg	gcctccacct	cctcctgggc	cgccctctac	ttcgtggccc	3240
tcattgacctt	cggcaactat	gtgctcttca	acctgctggg	ggccatcctc	gtggagggct	3300
tccaggcgga	gggcatgcc	aacagatccg	acacggacga	ggacaagacg	tcgggtccact	3360
tcgaggagga	cttccacaag	ctcagagaac	tccagaccac	agagctgaag	atgtgttccc	3420
tggccgtgac	ccccaacggg	cacctggagg	gacgaggcag	cctgtccctt	ccccctcatca	3480
atgcacacgt	tggcagcgcc	atgcctaccc	ccaagcagctc	accattcctg	gatgcagccc	3540
ccagcctccc	agactctcgg	cgtggcagca	gcagctccgg	ggaccgcga	ctgggagacc	3600
agaagcctcc	ggccagcctc	cgaagtcttc	cctgtgcccc	ctggggcccc	agtggcgccct	3660
ggagcagccg	gcgctccagc	tggagcagcc	tggggcgtgc	ccccagcctc	aagcgcccg	3720
gccagtgtgg	ggcaacgtgag	tccctgctgt	ctggcgaggg	caagggcagc	accgacgacg	3780
aagctgagga	cggcagggcc	gcgcccgggc	cccgtgccac	cccactgccc	cgggcccaggt	3840
ccctggaccc	acggccctctg	cggccggccg	ccctcccggc	taccaagtgc	cgcgatcgcg	3900
acgggcaggt	ggtggccctg	cccagcgact	tcttctctgcg	catcgacagc	caccgtgagg	3960
atgcacccga	gcttgacgac	gactcggagg	acagctgctg	cctccgcttg	cataaagtgc	4020
tggagcccta	caagccccag	tgggtccgga	gccgcgaggc	ctggggccctc	tacctcttct	4080
ccccacagaa	ccggttccgc	gtctctctgc	agaaggctcat	cacacacaag	atgtttgatc	4140
acgtggctct	cgtcttcac	ttcttcaact	gcgtcaccat	cgccctggag	aggcctgaca	4200
ttgacctcgg	cgaaccgag	cgggtcttcc	gggtcatcag	caattacatc	ttcacggcca	4260
tcttcgtggc	ggagatgatg	gtgaagggtg	tggccctggg	gtgctgtccc	ggcgagcacg	4320
cctacctgca	gagcagctgg	aacctgctgg	atgggctgct	ggtgctgggtg	tccctgggtgg	4380
acattgtcgt	ggccattggc	tcggctgggtg	gcgccaaagt	cctgggtggt	ctgcgcgtgc	4440
tggtggtgga	gacgtgata	tcgtcgtcca	ggcccatagg	gaacatcgtc	ctcatctgct	4500
gcgccttctt	catcattttt	ggcatcttgg	gtgtgcagct	cttcaaaggg	aagtctctact	4560
actgcgaggg	ccccgacacc	aggaacatct	ccaccaaggc	acagtgcggg	gccgcccact	4620
accgctgggt	gcgaacgcaag	tacaacttcg	tcaactctgg	ccaggccctg	atgtcgtgtg	4680
tcgtgctgtc	atccaaggat	ggatgggtga	acatcatgta	cgacgggctg	gatgccgtgg	4740
gtgtcgacca	gcagcctgtg	cagaaccaca	acccttggat	gctgctgtac	ttcatctcct	4800
tctgtctcat	cgtcagcttc	ttcgtgtcca	acatgttcgt	gggcgtcgtg	gtcgagaact	4860
tcacaagtgt	cgggcagcac	caggaggcgg	agggagcgcg	gcggcgagag	gagaagcggc	4920
tgccggcgct	agagaggagg	cgcaggagca	ctttccccag	cccagaggcc	cagcgccggc	4980
cctactatgc	cgactactcg	cccacgcgcc	gttccattca	ctcgtgtgtc	accagccact	5040
atctcgacct	cttcacaccc	ttcatcatct	gtgtcaacgt	catcaccatg	tccatggagc	5100
actataacca	acccaagtgc	ctggacgagg	ccctcaagta	ctgcaactac	gtcttcacca	5160
tcgtgtttgt	cttcgaggct	gcactgaagc	tggtagcatt	tgggttccgt	cggttcttca	5220
aggacaggtg	gaaccagctg	gacctggcca	tcgtgctgct	gtcactcatg	ggcatcacgc	5280
tggaggagat	agagatgagc	gccgcgctgc	ccatcaacc	caccatcatc	cgcacatgac	5340
gcgtgcttcg	cattgcccgt	gtgctgaagc	tgctgaagat	ggctacgggc	atgcgcgcc	5400
tgttggacac	tgtggtgcaa	gctctcccc	aggtggggaa	cctgggcctt	cttttcatgc	5460
tctgtttttt	tatctatgct	gcgctgggag	tggagctggt	cgggaggctg	gagtgcagtg	5520
aagacaaccc	ctgcgagggc	ctgagcaggc	acgccacctt	cagcaacttc	ggcatggcct	5580
tcctcacgct	gttccgcgtg	tccacggggg	acaactggaa	cgggatcatg	aaggacacgc	5640
tgcgcgagtg	ctcccgtgag	gacaagcact	gcctgagcta	cctgcgggcc	ctgtcgcccc	5700
tctacttcgt	gaccttcgtg	ctggtggccc	agttcgtgct	ggtgaacgtg	gtggtggccg	5760
tgctcatgaa	gcacctggag	gagagcaaca	aggaggcacg	ggaggatgcg	gagctggacg	5820
ccgagatcga	gctggagatg	gcgcaggggc	ccgggagtg	acgcccgggtg	gacgcggaca	5880
ggcctccctt	gccccaggag	agtcggggcg	caggggatgc	cccaaacctg	gttgacagca	5940
aggtgtccgt	gtccaggatg	ctctcgctgc	ccaacgacag	ctacatgttc	aggcccgtgg	6000
tgctgcctc	ggcgccccac	ccccggccgc	tgcaggaggt	ggagatggag	acctatgggg	6060
ccggcacccc	cttgggctcc	gttgccctct	tgcactctcc	gcccgcagag	tcctgtgcct	6120
ccctccagat	cccactgggt	gtgtcgctcc	cagccaggag	cggcgagccc	ctccagccc	6180
tgctccctcg	gggcacagcc	cgctccccc	gtctcagccg	gctgctctgc	agacaggagg	6240
ctgtgcacac	cgattccttg	gaagggaaga	ttgacagccc	tagggacacc	ctggatcctg	6300
						6360

```

cagagcctgg tgagaaaacc cgggtgagggc cgggtgaccca ggggggctcc ctgcagtccc 6420
caccacgctc cccacggccc gccagcgctc gcactcgtaa gcataccttc ggacagcact 6480
gcgtctccag ccggccggcg gccccaggcg gagaggaggc cgaggcctcg gaccagccg 6540
acgaggaggt cagccacatc accagctccg cctgcccctg gcagcccaca gccgagcccc 6600
atggccccga agcctctccg gtggccggcg gcgagcggga cctgcgcagg ctctacagcg 6660
tggagctca gggcttctcg gacaagccgg gccgggcaga cgagcagtgg cggccctcgg 6720
cggagctggg cagcggggag cctggggagg cgaagccctg gggccctgag gccgagcccc 6780
ctctgggtgc gcgcagaaag aagaagatga gccccccctg catctcgggtg gaacccctcg 6840
cggaggacga gggctctgcg cggccctccg cggcagaggg cggcagcacc aactgaggc 6900
gcaggacccc gtcctgtgag gccacgcctc acagggactc cctggagccc acagagggt 6960
caggcgccgg gggggaccct gcagccaagg gggagcgcgt gggccaggcc tcctgccggg 7020
ctgagcacct gaccgtcccc agctttgcct ttgagccgct ggacctcggg gtccccagtg 7080
gagacccttt cttggacggg agccacagtg tgaccccaga atccagagct tcctcttcag 7140
gggccatagt gcccttgaa ccccagaat cagagcctcc catgcccgtc ggtgaccccc 7200
cagagaagag gcgggggctg tacctcacag tccccagtg tcctctggag aaaccagggt 7260
ccccctcagc caccctgcc ccagggggtg gtgcagatga ccccgtag ctcggggctt 7320
ggtgccgccc acggctttgg ccctggggtc tgggggcccc gctggggtag aggccaggc 7380
agaacctgc atggaccctg acttgggtcc cgctgtgagc agaaaggccc ggggaggatg 7440
acggcccagg ccctgggtct ctgccagcg aagcaggagt agctgccggg ccccacgagc 7500
ctccatccgt tctggttcgg gtttctccga gtttctgtac cagccgaggc tgtgcgggca 7560
actgggtcag cctcccgtea ggagagaagc cgctctgtg ggacgaagac cgggcacccc 7620
ccagagaggg gaaggtacca ggttgcgtcc ttccaggccc cgctgtgta caggacactc 7680
gctggggggc ctgtgccctt gccggcggca ggttcagcc accgcggccc aatgtcacct 7740
tcactcacag tctgagttct tgtccgcctg tcacgcctc accaccctcc cttccagcc 7800
accacccttt ccgttccgct cgggccttc cagaagcgtc ctgtgactct gggagagggtg 7860
acacctcact aaggggccga ccccatggag taacgcgc 7898

```

<210> 4
 <211> 2353
 <212> PRT
 <213> Homo Sapiens

```

<400> 4
Met Thr Glu Gly Ala Arg Ala Ala Asp Glu Val Arg Val Pro Leu Gly
 1          5          10          15
Ala Pro Pro Pro Gly Pro Ala Ala Leu Val Gly Ala Ser Pro Glu Ser
 20          25          30
Pro Gly Ala Pro Gly Arg Glu Ala Glu Arg Gly Ser Glu Leu Gly Val
 35          40          45
Ser Pro Ser Glu Ser Pro Ala Ala Glu Arg Gly Ala Glu Leu Gly Ala
 50          55          60
Asp Glu Glu Gln Arg Val Pro Tyr Pro Ala Leu Ala Ala Thr Val Phe
 65          70          75          80
Phe Cys Leu Gly Gln Thr Thr Arg Pro Arg Ser Trp Cys Leu Arg Leu
 85          90          95
Val Cys Asn Pro Trp Phe Glu His Val Ser Met Leu Val Ile Met Leu
100          105          110
Asn Cys Val Thr Leu Gly Met Phe Arg Pro Cys Glu Asp Val Glu Cys
115          120          125
Gly Ser Glu Arg Cys Asn Ile Leu Glu Ala Phe Asp Ala Phe Ile Phe
130          135          140
Ala Phe Phe Ala Val Glu Met Val Ile Lys Met Val Ala Leu Gly Leu
145          150          155          160
Phe Gly Gln Lys Cys Tyr Leu Gly Asp Thr Trp Asn Arg Leu Asp Phe
165          170          175
Phe Ile Val Val Ala Gly Met Met Glu Tyr Ser Leu Asp Gly His Asn
180          185          190
Val Ser Leu Ser Ala Ile Arg Thr Val Arg Val Leu Arg Pro Leu Arg
195          200          205
Ala Ile Asn Arg Val Pro Ser Met Arg Ile Leu Val Thr Leu Leu Leu

```

210	215	220
Asp Thr Leu Pro Met Leu	Gly Asn Val Leu Leu	Leu Cys Phe Phe Val
225	230	235
Phe Phe Ile Phe Gly Ile	Val Gly Val Gln Leu	Trp Ala Gly Leu Leu
245	250	255
Arg Asn Arg Cys Phe Leu	Asp Ser Ala Phe Val	Arg Asn Asn Asn Leu
260	265	270
Thr Phe Leu Arg Pro Tyr	Tyr Gln Thr Glu Glu Gly	Glu Glu Asn Pro
275	280	285
Phe Ile Cys Ser Ser Arg	Arg Asp Asn Gly Met Gln	Lys Cys Ser His
290	295	300
Ile Pro Gly Arg Arg Glu	Leu Arg Met Pro Cys Thr	Leu Gly Trp Glu
305	310	315
Ala Tyr Thr Gln Pro Gln	Ala Glu Gly Val Gly Ala	Ala Arg Asn Ala
325	330	335
Cys Ile Asn Trp Asn Gln	Tyr Tyr Asn Val Cys Arg	Ser Gly Asp Ser
340	345	350
Asn Pro His Asn Gly Ala	Ile Asn Phe Asp Asn Ile	Gly Tyr Ala Trp
355	360	365
Ile Ala Ile Phe Gln Val	Ile Thr Leu Glu Gly Trp	Val Asp Ile Met
370	375	380
Tyr Tyr Val Met Asp Ala	His Ser Phe Tyr Asn Phe	Ile Tyr Phe Ile
385	390	395
Leu Leu Ile Ile Val Gly	Ser Phe Phe Met Ile Asn	Leu Cys Leu Val
405	410	415
Val Ile Ala Thr Gln Phe	Ser Glu Thr Lys Gln Arg	Glu Ser Gln Leu
420	425	430
Met Arg Glu Gln Arg Ala	Arg His Leu Ser Asn Asp	Ser Thr Leu Ala
435	440	445
Ser Phe Ser Glu Pro Gly	Ser Cys Tyr Glu Glu Leu	Leu Lys Tyr Val
450	455	460
Gly His Ile Phe Arg Lys	Val Lys Arg Arg Ser Leu	Arg Leu Tyr Ala
465	470	475
Arg Trp Gln Ser Arg Trp	Arg Lys Lys Val Asp Pro	Ser Ala Val Gln
485	490	495
Gly Gln Gly Pro Gly His	Arg Gln Arg Ala Gly Arg	His Thr Ala
500	505	510
Ser Val His His Leu Val	Tyr His His His His His	His His His His
515	520	525
Tyr His Phe Ser His Gly	Ser Pro Arg Arg Pro Gly	Pro Glu Pro Gly
530	535	540
Ala Cys Asp Thr Arg Leu	Val Arg Ala Gly Ala Pro	Pro Ser Pro Pro
545	550	555
Ser Pro Gly Arg Gly Pro	Pro Asp Ala Glu Ser Val	His Ser Ile Tyr
565	570	575
His Ala Asp Cys His Ile	Glu Gly Pro Gln Glu Arg	Ala Arg Val Ala
580	585	590
His Ala Ala Ala Thr Ala	Ala Ala Ser Leu Arg Leu	Ala Thr Gly Leu
595	600	605
Gly Thr Met Asn Tyr Pro	Thr Ile Leu Pro Ser Gly	Val Gly Ser Gly
610	615	620
Lys Gly Ser Thr Ser Pro	Gly Pro Lys Gly Lys Trp	Ala Gly Gly Pro
625	630	635
Pro Gly Thr Gly Gly His	Gly Pro Leu Ser Leu Asn	Ser Pro Asp Pro
645	650	655
Tyr Glu Lys Ile Pro His	Val Val Gly Glu His Gly	Leu Gly Gln Ala
660	665	670
Pro Gly His Leu Ser Gly	Leu Ser Val Pro Cys Pro	Leu Pro Ser Pro
675	680	685
Pro Ala Gly Thr Leu Thr	Cys Glu Leu Lys Ser Cys	Pro Tyr Cys Thr
690	695	700
Arg Ala Leu Glu Asp Pro	Glu Gly Glu Leu Ser Gly	Ser Glu Ser Gly

705					710					715				720
Asp	Ser	Asp	Gly	Arg	Gly	Val	Tyr	Glu	Phe	Thr	Gln	Asp	Val	Arg
				725					730					735
Gly	Asp	Arg	Trp	Asp	Pro	Thr	Arg	Pro	Pro	Arg	Ala	Thr	Asp	Thr
			740					745					750	
Gly	Pro	Gly	Pro	Gly	Ser	Pro	Gln	Arg	Arg	Ala	Gln	Gln	Arg	Ala
		755					760					765		Ala
Pro	Gly	Glu	Pro	Gly	Trp	Met	Gly	Arg	Leu	Trp	Val	Thr	Phe	Ser
	770					775					780			Gly
Lys	Leu	Arg	Arg	Ile	Val	Asp	Ser	Lys	Tyr	Phe	Ser	Arg	Gly	Ile
785					790					795				Met
Met	Ala	Ile	Leu	Val	Asn	Thr	Leu	Ser	Met	Gly	Val	Glu	Tyr	His
				805					810					815
Gln	Pro	Glu	Glu	Leu	Thr	Asn	Ala	Leu	Glu	Ile	Ser	Asn	Ile	Val
			820					825					830	Phe
Thr	Ser	Met	Phe	Ala	Leu	Glu	Met	Leu	Leu	Lys	Leu	Leu	Ala	Cys
		835					840					845		Gly
Pro	Leu	Gly	Tyr	Ile	Arg	Asn	Pro	Tyr	Asn	Ile	Phe	Asp	Gly	Ile
	850					855					860			Ile
Val	Val	Ile	Ser	Val	Trp	Glu	Ile	Val	Gly	Gln	Ala	Asp	Gly	Gly
865					870				875					Leu
Ser	Val	Leu	Arg	Thr	Phe	Arg	Leu	Leu	Arg	Val	Leu	Lys	Leu	Val
				885					890					Arg
Phe	Leu	Pro	Ala	Leu	Arg	Arg	Gln	Leu	Val	Val	Leu	Val	Lys	Thr
			900					905					910	Met
Asp	Asn	Val	Ala	Thr	Phe	Cys	Thr	Leu	Leu	Met	Leu	Phe	Ile	Phe
	915						920					925		Ile
Phe	Ser	Ile	Leu	Gly	Met	His	Leu	Phe	Gly	Cys	Lys	Phe	Ser	Leu
	930				935						940			Lys
Thr	Asp	Thr	Gly	Asp	Thr	Val	Pro	Asp	Arg	Lys	Asn	Phe	Asp	Ser
945					950					955				Leu
Leu	Trp	Ala	Ile	Val	Thr	Val	Phe	Gln	Ile	Leu	Thr	Gln	Glu	Asp
			965						970					Trp
Asn	Val	Val	Leu	Tyr	Asn	Gly	Met	Ala	Ser	Thr	Ser	Ser	Trp	Ala
			980				985						990	Ala
Leu	Tyr	Phe	Val	Ala	Leu	Met	Thr	Phe	Gly	Asn	Tyr	Val	Leu	Phe
	995					1000						1005		Asn
Leu	Leu	Val	Ala	Ile	Leu	Val	Glu	Gly	Phe	Gln	Ala	Glu	Gly	Asp
	1010					1015					1020			Ala
Asn	Arg	Ser	Asp	Thr	Asp	Glu	Asp	Lys	Thr	Ser	Val	His	Phe	Glu
1025					1030					1035				Glu
Asp	Phe	His	Lys	Leu	Arg	Glu	Leu	Gln	Thr	Thr	Glu	Leu	Lys	Met
			1045						1050					Cys
Ser	Leu	Ala	Val	Thr	Pro	Asn	Gly	His	Leu	Glu	Gly	Arg	Gly	Ser
			1060					1065					1070	Leu
Ser	Pro	Pro	Leu	Ile	Met	Cys	Thr	Ala	Ala	Thr	Pro	Met	Pro	Thr
	1075					1080					1085			Pro
Lys	Ser	Ser	Pro	Phe	Leu	Asp	Ala	Ala	Pro	Ser	Leu	Pro	Asp	Ser
	1090					1095					1100			Arg
Arg	Gly	Ser	Ser	Ser	Ser	Gly	Asp	Pro	Pro	Leu	Gly	Asp	Gln	Lys
1105					1110					1115				Pro
Pro	Ala	Ser	Leu	Arg	Ser	Ser	Pro	Cys	Ala	Pro	Trp	Gly	Pro	Ser
			1125						1130				1135	Gly
Ala	Trp	Ser	Ser	Arg	Arg	Ser	Ser	Trp	Ser	Ser	Leu	Gly	Arg	Ala
			1140					1145					1150	Pro
Ser	Leu	Lys	Arg	Arg	Gly	Gln	Cys	Gly	Glu	Arg	Glu	Ser	Leu	Leu
	1155					1160					1165			Ser
Gly	Glu	Gly	Lys	Gly	Ser	Thr	Asp	Asp	Glu	Ala	Glu	Asp	Gly	Arg
	1170					1175					1180			Ala
Ala	Pro	Gly	Pro	Arg	Ala	Thr	Pro	Leu	Arg	Arg	Ala	Glu	Ser	Leu
1185					1190					1195				Asp
Pro	Arg	Pro	Leu	Arg	Pro	Ala	Ala	Leu	Pro	Pro	Thr	Lys	Cys	Arg

				1205						1210						1215		
Arg	Asp	Gly	Gln	Val	Val	Ala	Leu	Pro	Ser	Asp	Phe	Phe	Leu	Arg	Ile			
			1220					1225					1230					
Asp	Ser	His	Arg	Glu	Asp	Ala	Ala	Glu	Leu	Asp	Asp	Asp	Ser	Glu	Asp			
		1235					1240					1245						
Ser	Cys	Cys	Leu	Arg	Leu	His	Lys	Val	Leu	Glu	Pro	Tyr	Lys	Pro	Gln			
	1250					1255					1260							
Trp	Cys	Arg	Ser	Arg	Glu	Ala	Trp	Ala	Leu	Tyr	Leu	Phe	Ser	Pro	Gln			
1265					1270					1275					1280			
Asn	Arg	Phe	Arg	Val	Ser	Cys	Gln	Lys	Val	Ile	Thr	His	Lys	Met	Phe			
				1285					1290					1295				
Asp	His	Val	Val	Leu	Val	Phe	Ile	Phe	Leu	Asn	Cys	Val	Thr	Ile	Ala			
			1300					1305					1310					
Leu	Glu	Arg	Pro	Asp	Ile	Asp	Pro	Gly	Ser	Thr	Glu	Arg	Val	Phe	Leu			
		1315					1320				1325							
Ser	Val	Ser	Asn	Tyr	Ile	Phe	Thr	Ala	Ile	Phe	Val	Ala	Glu	Met	Met			
	1330					1335					1340							
Val	Lys	Val	Val	Ala	Leu	Gly	Leu	Leu	Ser	Gly	Glu	His	Ala	Tyr	Leu			
1345				1350						1355					1360			
Gln	Ser	Ser	Trp	Asn	Leu	Leu	Asp	Gly	Leu	Leu	Val	Leu	Val	Ser	Leu			
				1365					1370					1375				
Val	Asp	Ile	Val	Val	Ala	Met	Ala	Ser	Ala	Gly	Gly	Ala	Lys	Ile	Leu			
			1380					1385					1390					
Gly	Val	Leu	Arg	Val	Leu	Arg	Leu	Leu	Arg	Thr	Leu	Arg	Pro	Leu	Arg			
	1395						1400				1405							
Val	Ile	Ser	Arg	Ala	Pro	Gly	Leu	Lys	Leu	Val	Val	Glu	Thr	Leu	Ile			
	1410					1415				1420								
Ser	Ser	Leu	Arg	Pro	Ile	Gly	Asn	Ile	Val	Leu	Ile	Cys	Cys	Ala	Phe			
1425				1430					1435					1440				
Phe	Ile	Ile	Phe	Gly	Ile	Leu	Gly	Val	Gln	Leu	Phe	Lys	Gly	Lys	Phe			
				1445				1450						1455				
Tyr	Tyr	Cys	Glu	Gly	Pro	Asp	Thr	Arg	Asn	Ile	Ser	Thr	Lys	Ala	Gln			
		1460						1465				1470						
Cys	Arg	Ala	Ala	His	Tyr	Arg	Trp	Val	Arg	Arg	Lys	Tyr	Asn	Phe	Asp			
	1475						1480				1485							
Asn	Leu	Gly	Gln	Ala	Leu	Met	Ser	Leu	Phe	Val	Leu	Ser	Ser	Lys	Asp			
	1490					1495				1500								
Gly	Trp	Val	Asn	Ile	Met	Tyr	Asp	Gly	Leu	Asp	Ala	Val	Gly	Val	Asp			
1505				1510					1515									

1700					1705					1710					
Ile	Ile	Arg	Ile	Met	Arg	Val	Leu	Arg	Ile	Ala	Arg	Val	Leu	Lys	Leu
		1715					1720					1725			
Leu	Lys	Met	Ala	Thr	Gly	Met	Arg	Ala	Leu	Leu	Asp	Thr	Val	Val	Gln
		1730					1735					1740			
Ala	Leu	Pro	Gln	Val	Gly	Asn	Leu	Gly	Leu	Leu	Phe	Met	Leu	Leu	Phe
		1745					1750					1755			
Phe	Ile	Tyr	Ala	Ala	Leu	Gly	Val	Glu	Leu	Phe	Gly	Arg	Leu	Glu	Cys
				1765					1770					1775	
Ser	Glu	Asp	Asn	Pro	Cys	Glu	Gly	Leu	Ser	Arg	His	Ala	Thr	Phe	Ser
			1780						1785					1790	
Asn	Phe	Gly	Met	Ala	Phe	Leu	Thr	Leu	Phe	Arg	Val	Ser	Thr	Gly	Asp
			1795						1800					1805	
Asn	Trp	Asn	Gly	Ile	Met	Lys	Asp	Thr	Leu	Arg	Glu	Cys	Ser	Arg	Glu
			1810						1815					1820	
Asp	Lys	His	Cys	Leu	Ser	Tyr	Leu	Pro	Ala	Leu	Ser	Pro	Val	Tyr	Phe
			1825						1830					1835	
Val	Thr	Phe	Val	Leu	Val	Ala	Gln	Phe	Val	Leu	Val	Asn	Val	Val	Val
				1845					1850					1855	
Ala	Val	Leu	Met	Lys	His	Leu	Glu	Glu	Ser	Asn	Lys	Glu	Ala	Arg	Glu
			1860						1865					1870	
Asp	Ala	Glu	Leu	Asp	Ala	Glu	Ile	Glu	Leu	Glu	Met	Ala	Gln	Gly	Pro
			1875						1880					1885	
Gly	Ser	Ala	Arg	Arg	Val	Asp	Ala	Asp	Arg	Pro	Pro	Leu	Pro	Gln	Glu
			1890						1895					1900	
Ser	Pro	Gly	Ala	Arg	Asp	Ala	Pro	Asn	Leu	Val	Ala	Arg	Lys	Val	Ser
			1905						1910					1915	
Val	Ser	Arg	Met	Leu	Ser	Leu	Pro	Asn	Asp	Ser	Tyr	Met	Phe	Arg	Pro
				1925					1930					1935	
Val	Val	Pro	Ala	Ser	Ala	Pro	His	Pro	Arg	Pro	Leu	Gln	Glu	Val	Glu
			1940						1945					1950	
Met	Glu	Thr	Tyr	Gly	Ala	Gly	Thr	Pro	Leu	Gly	Ser	Val	Ala	Ser	Val
			1955						1960					1965	
His	Ser	Pro	Pro	Ala	Glu	Ser	Cys	Ala	Ser	Leu	Gln	Ile	Pro	Leu	Ala
			1970						1975					1980	
Val	Ser	Ser	Pro	Ala	Arg	Ser	Gly	Glu	Pro	Leu	His	Ala	Leu	Ser	Pro
			1985						1990					1995	
Arg	Gly	Thr	Ala	Arg	Ser	Pro	Ser	Leu	Ser	Arg	Leu	Leu	Cys	Arg	Gln
				2005					2010					2015	
Glu	Ala	Val	His	Thr	Asp	Ser	Leu	Glu	Gly	Lys	Ile	Asp	Ser	Pro	Arg
			2020						2025					2030	
Asp	Thr	Leu	Asp	Pro	Ala	Glu	Pro	Gly	Glu	Lys	Thr	Pro	Val	Arg	Pro
			2035						2040					2045	
Val	Thr	Gln	Gly	Gly	Ser	Leu	Gln	Ser	Pro	Pro	Arg	Ser	Pro	Arg	Pro
			2050						2055					2060	
Ala	Ser	Val	Arg	Thr	Arg	Lys	His	Thr	Phe	Gly	Gln	His	Cys	Val	Ser
			2065						2070					2075	
Ser	Arg	Pro	Ala	Ala	Pro	Gly	Gly	Glu	Glu	Ala	Glu	Ala	Ser	Asp	Pro
				2085					2090					2095	
Ala	Asp	Glu	Glu	Val	Ser	His	Ile	Thr	Ser	Ser	Ala	Cys	Pro	Trp	Gln
			2100						2105					2110	
Pro	Thr	Ala	Glu	Pro	His	Gly	Pro	Glu	Ala	Ser	Pro	Val	Ala	Gly	Gly
			2115						2120					2125	
Glu	Arg	Asp	Leu	Arg	Arg	Leu	Tyr	Ser	Val	Asp	Ala	Gln	Gly	Phe	Leu
			2130						2135					2140	
Asp	Lys	Pro	Gly	Arg	Ala	Asp	Glu	Gln	Trp	Arg	Pro	Ser	Ala	Glu	Leu
			2145						2150					2155	
Gly	Ser	Gly	Glu	Pro	Gly	Glu	Ala	Lys	Ala	Trp	Gly	Pro	Glu	Ala	Glu
				2165					2170					2175	
Pro	Ala	Leu	Gly	Ala	Arg	Arg	Lys	Lys	Lys	Met	Ser	Pro	Pro	Cys	Ile
			2180						2185					2190	
Ser	Val	Glu	Pro	Pro	Ala	Glu	Asp	Glu	Gly	Ser	Ala	Arg	Pro	Ser	Ala

2195										2200					2205				
Ala	Glu	Gly	Gly	Ser	Thr	Thr	Leu	Arg	Arg	Arg	Thr	Pro	Ser	Cys	Glu				
2210										2215					2220				
Ala	Thr	Pro	His	Arg	Asp	Ser	Leu	Glu	Pro	Thr	Glu	Gly	Ser	Gly	Ala				
2225										2230					2235				
Gly	Gly	Asp	Pro	Ala	Ala	Lys	Gly	Glu	Arg	Trp	Gly	Gln	Ala	Ser	Cys				
2245										2250					2255				
Arg	Ala	Glu	His	Leu	Thr	Val	Pro	Ser	Phe	Ala	Phe	Glu	Pro	Leu	Asp				
2260										2265					2270				
Leu	Gly	Val	Pro	Ser	Gly	Asp	Pro	Phe	Leu	Asp	Gly	Ser	His	Ser	Val				
2275										2280					2285				
Thr	Pro	Glu	Ser	Arg	Ala	Ser	Ser	Ser	Gly	Ala	Ile	Val	Pro	Leu	Glu				
2290										2295					2300				
Pro	Pro	Glu	Ser	Glu	Pro	Pro	Met	Pro	Val	Gly	Asp	Pro	Pro	Glu	Lys				
2305										2310					2315				
Arg	Arg	Gly	Leu	Tyr	Leu	Thr	Val	Pro	Gln	Cys	Pro	Leu	Glu	Lys	Pro				
2325										2330					2335				
Gly	Ser	Pro	Ser	Ala	Thr	Pro	Ala	Pro	Gly	Gly	Gly	Ala	Asp	Asp	Pro				
2340										2345					2350				
Val																			

<210> 5
 <211> 7364.
 <212> DNA
 <213> Homo Sapiens

<400> 5

gcgggcgcggt	ctggcgcggt	ggggcggggc	gaggtccgct	gcggtcccg	cggtccggt	60
gctgctccgc	tctgagcgcc	tggcgcgccc	cgcgccctcc	ctggcggggc	cgctggggcg	120
gggatgcacg	cggggcccgg	gagccatggt	ccgcttcggg	gacgagctgg	gcggcgcgta	180
tggaggcccc	ggcgggcgag	agcgggcccc	ggcgggcggg	gccggcgggg	cgggggggccc	240
gggtcccggt	gggctgcagc	ccggccagcg	ggtcctctac	aagcaatcga	tcgcgcagcg	300
cgcgcgacc	atggcgctgt	acaaccccat	cccggtcaag	cagaactgct	tcaccgtcaa	360
ccgctcgctc	ttcgtcttca	gcgaggacaa	cgctgtccgc	aaatacgcg	agcgcatcac	420
cgagtggcct	ccattcgagt	atatgatcct	ggccaccatc	atcgccaact	gcatcggtgt	480
ggccctggag	cagcacctcc	ctgatgggga	caaaacgccc	atgtccgagc	ggctggacga	540
cacggagccc	tatttcacg	ggatcttttg	cttcgaggca	gggatcaaaa	tcacgtctct	600
gggctttgtc	ttccacaagg	gctcttacct	gcggaacggc	tggaaactga	tggacttcgt	660
ggtcgctctc	acagggatcc	ttgccacggc	tggaaactga	ttcgacctgc	gaacactgag	720
ggctgtgctg	gtgctgaggg	ccctgaagct	ggtgtctggg	attccaagtt	tgcaggtggt	780
gctcaagtcc	atcatgaagg	ccatgggtcc	actcctcgag	attgggctgc	ttctcttctt	840
tgccatcctc	atgtttgcca	tcattggcct	ggagttctac	atgggcaagt	tccacaaggc	900
ctgtttcccc	aacagcacag	atgcgagacc	cgtgggtgac	ttcccctgtg	gcaaggaggc	960
cccagcccgg	ctgtgcgagg	gcgacactga	gtgcccggag	tactggccag	gacccaactt	1020
tggcatcacc	aactttgaca	atatacctgt	tgccatcttg	acggtgttcc	agtgcatacc	1080
catggagggc	tggactgaca	tcctctataa	tacaaacgat	gcggccggca	acacctggaa	1140
ctggctctac	ttcatccctc	tcatacatcat	cggctccttc	ttcatgtctc	acctggtgct	1200
gggctgtctc	tcgggggagt	ttgccaagga	gcgagagagg	gtggagaacc	gccgcgcctt	1260
cctgaagctg	cgccggcgag	agcagatcga	gcgagagctc	aacgggtacc	tggagtggat	1320
cttcaaggcg	gaggaagtca	tgctggccga	ggaggacagg	aatgcagagg	agaagtcccc	1380
tttgacgtg	ctgaagagag	cggccaccaa	gaagagcaga	aatgacctga	tccacgcaga	1440
ggaggagag	gaccggtttg	cagatctctg	tgctgttgga	tcccccttcg	cccgcgccag	1500
cctcaagagc	gggaagacag	agagctcgtc	atacttccgg	aggaaggaga	agatgttccg	1560
gttttttatc	cggcgcatgg	tgaaggctca	gagcttctac	tgggtgtgtg	tggtgtgtgt	1620
ggcctgaac	acactgtgtg	tggccatggt	gcattacaac	cagccgcggc	ggcttaccac	1680
gacctgtat	tttgagagt	ttgttttcct	gggtctcttc	ctcacagaga	tgtccctgaa	1740
gatgtatggc	ctggggccca	gaagctactt	ccggtctctc	ttcaactgct	tcgactttgg	1800
ggtcactcgt	ctggagcgtc	ttgaagtgg	ctggcgcgcc	atcaagccgg	gaagctcctt	1860
tgggatcagt	gtgctgcggg	ccctccgcct	gctgaggatc	ttcaaagtca	cgaagtactg	1920
gagctccctg	cggaaacctg	tggtgtccct	gctgaactcc	atgaagtcca	tcatacagct	1980
gctcttcttg	ctcttctctg	tcattgtgg	cttcgccttg	ctggggatgc	agctgtttgg	2040

gggacagttc	aacttccagg	atgagactcc	cacaaccaac	ttcgacacct	tccctgcccg	2100
catcctcaact	gtcttccaga	tcctgacggg	agaggactgg	aatgcagtga	tgtatcacgg	2160
gatcgaatcg	caaggcgggc	tcagcaaagg	catgtttctg	tccttttact	tcattgtcct	2220
gacactgttc	ggaaactaca	ctctgctgaa	tgtctttctg	gccatcgctg	tggacaacct	2280
ggccaacgcc	caagagctga	ccaaggatga	agaggagatg	gaagaagcag	ccaatcagaa	2340
gcttgctctg	caaaaaggcca	aagaagtggc	tgaagtcagc	cccatgtctg	ccgcgaacat	2400
ctccatcgcc	gccaggcagc	agaactcggc	caaggcgcg	tcggtgtggg	agcagcgggc	2460
cagccagcta	cggctgcaga	acctgcgggc	cagctgcgag	gcgctgtaca	gcgagatgga	2520
ccccgaggag	cggctgcgct	tcgccactac	gcgccacctg	cggcccgaca	tgaagacgca	2580
cctggaccgg	cgcgtggtgg	tggagctggg	ccgcgacggc	gcgcgggggc	ccgtgggagg	2640
caaaagcccg	cctgaggctg	cggaggcccc	cggggcgctc	gacctccgc	gcaggcacca	2700
ccggcaccgc	gacaaggaca	agacccccgc	ggcgggggac	caggaccgag	cagaggcccc	2760
gaaggcgagg	agcgggggag	ccggtgcccc	ggaggagcgg	ccgcggccgc	accgcagcca	2820
cagcaaggag	gccgcggggc	ccccggaggc	gcggagcgag	cgcgcccgag	gcccaggccc	2880
cgagggcgcc	cggcgccacc	accggcgcg	ctccccggag	gaggcgcccg	agcgggagcc	2940
ccgacgccac	cgcgcgcacc	ggcaccagga	tccgagcaag	gagtgcgccc	gcgccaaggg	3000
cgagcgggcg	gcgcggcacc	gcggcgggcc	ccgagcgggg	ccccgggagg	cggagagcgg	3060
ggaggagccg	gcgcggcgcc	accggggccc	gcacaaggcg	cagcctgctc	acgaggctgt	3120
ggagaaggag	accacggaga	aggaggccac	ggaggaaggag	gctgagatag	tggagaccga	3180
caaggaaaag	gagctccgga	accaccagcc	ccgggagcca	cactgtgacc	tggagaccag	3240
tgggactgtg	actgtgggtc	ccatgcacac	actgcccagc	acctgtctcc	agaagggtga	3300
ggaacagcca	gaggatgcag	acaatcacgg	gaacgtcact	cgcatgggca	gtcagcccc	3360
agaccggaac	actattgtac	atatcccagt	gagctgtacg	ggccctcttg	gggaaggccac	3420
ggtcgttccc	agtggtaacg	tggacctgga	aagccaagca	gaggggaaga	aggagggtga	3480
agcggatgac	gtgatgagga	gcggcccccg	gcctatcgte	ccatacagct	ccatgttctg	3540
tttaagcccc	accaaacctgc	tcgcgcgctt	ctgccactac	atcgtgacca	tgaggctact	3600
cgaggtggte	attctcgtgg	tcatacgctt	gagcagcatc	gccctggctg	ctgaggacc	3660
agtgcgcaca	gactcgcoca	ggaacaacgc	tctgaaatac	ctggattaca	ttttacttgg	3720
tgtctttacc	tttgagatgg	tgtataaagat	gatcgacttg	ggactgctgc	ttcaccctgg	3780
agcctatttc	cgggacttgt	ggaacatttc	ggacttcatt	gtggtcagtg	gcgccttggt	3840
ggcggttgct	ttctcaggat	ccaaagggaa	agacatcaat	accatcaagt	ctctgagagt	3900
ccttcgtgtc	ctgcggcccc	tcaagaccat	caaacggctg	cccaagctca	aggctgtggt	3960
tgaactgtgt	gtgaactccc	tgaagaatgt	cctcaacatc	ttgattgtct	acatgtctct	4020
catgttcata	tttgccgtca	ttgcgggtga	gctcttcaaa	gggaagtgtt	tctactgcac	4080
agatgcgaat	aaggagctgg	agaggggactg	cagggggtcag	tatttggatt	atgagaagga	4140
ggaagtggaa	gctcagccca	ggcagtgga	gaaatacgac	tttctactacg	acaatgtgct	4200
ctgggctctg	ctgacgctgt	tcacagtgtc	cacgggagaa	ggctggccca	tgggtgctgaa	4260
acactccgtg	gatgccacct	atgaggagca	gggtccaagc	cctgggtacc	gcatggagct	4320
gtccatcttc	tacgtggctc	actttgtggt	ctttcccttc	ttcttctgca	acatctttgt	4380
ggctttgatc	atcatcacct	tccaggagca	gggggacaag	gtgatgtctg	aatgcagcct	4440
ggagaagaac	gagagggctt	gcattgactt	cgccatcagc	gccaaacccc	tgacacggta	4500
catgccccaa	aaccggcagt	cgttccagta	taagacgtgg	acatttgtgg	tctccccgcc	4560
ctttgaatac	ttcatcatgg	ccatgatagc	cctcaacact	gtgggtgctga	tgatgaagtt	4620
ctatgatgca	ccctatgagt	acgagctgat	gctgaaatgc	ctgaacatcg	tgttcacatc	4680
catgtttctc	atgggaatgcg	tgtggaagat	catcgctttt	gggggtgctga	actattttcag	4740
agatgcctgg	aatgtctttg	actttgtcac	tgtgttggga	agtattactg	atatttttagt	4800
aacagagatt	gcggaaaacga	acaatttcat	caacctcagc	ttcctccgcc	tctttcgagc	4860
tgcgcggctg	atcaagctgc	tccgccaggg	ctacaccatc	cgcatcctgc	tgtggacctt	4920
tgtccagctc	ttcaaggccc	tgccctacgt	gtgtctgtct	attgccatgc	tgttcttcat	4980
ctacgccatc	atcggcatgc	aggtgttttg	gaatattgcc	ctggatgatg	acaccagcat	5040
caaccgccac	aacaacttcc	ggacgttttt	ctaagccctg	atgctgctgt	tcaggagcgc	5100
cacgggggag	gcctggcacg	agatcatgct	gtcctgcctg	agcaaccagg	cctgtgatga	5160
gcaggccaat	gccaccgagt	gtggaagtga	ctttgcctac	ttctacttcg	tctccttcat	5220
cttctctgtc	tccttttctga	tgttgaacct	ctttgtggct	gtgatcatgg	acaattttga	5280
gtaccctcag	cgccactctt	ccatcctagg	tccctaccac	ttggatgagt	tcacccgggt	5340
ctgggctgaa	tacgaccggg	ctgcgtgtgg	gcgcatacgt	tacaatgaca	tgtttgagat	5400
gctgaaacac	atgtccccgc	ctctggggct	gggggaagaaa	tgccctgtct	gagttgctta	5460
caagcgccctg	gttcgcatga	acatgcccac	ctccaacgag	gacatgactg	ttcactttcac	5520
gtccacgctg	atggccctca	tccggacggc	actggagatc	aagctggccc	cagctgggac	5580
aaagcagcat	cagtgtgacg	cggagtttag	gaaggagatt	tccgttgtgt	gggccaatct	5640
gccccagaag	actttggact	tgctgggtacc	accccataag	cctgatgaga	tgacagtggg	5700
gaagggttat	gcagctctga	tgatatttga	cttctacaag	cagaacaaaa	ccaccagaga	5760

```

ccagatgcag caggctcctg gaggcctctc ccagatgggt cctgtgtccc tgttccaccc 5820
tctgaaggcc accctggagc agacacagcc ggctgtgctc cgaggagccc gggttttcct 5880
tcgacagaag agttccacct ccctcagcaa tggcggggcc atacaaaacc aagagagtgg 5940
catcaaagag tctgtctcct ggggcactca aaggaccagc gatgcacccc atgaggccag 6000
gccacccctg gagcgtggcc actccacaga gatccctgtg gggcggtcag gagcactggc 6060
tgttggacgtt cagatgcaga gcataaccog gaggggccct gatggggagc cccagcctgg 6120
gctggagagc cagggtcgag cggcctccat gccccgcctt gcggccgaga ctcagcccg 6180
cacagatgcc agcccatga agcgtccat ctccacgctg gccagcggc cccgtgggac 6240
tcattcttgc agcaccaccc cggaccgccc acccctagc caggcgtcgt cgcaccacca 6300
ccaccaccgc tgcaccgcc gcagggacag gaagcagagg tccctggaga aggggccag 6360
cctgtctgcc gatatggatg gcgcaccaag cagtgtgtg gggccggggc tgcccccg 6420
agaggggccc acaggctgcc ggcgggaacg agagcgccgg caggagcggg gccggtccc 6480
ggagcggagg cagccctcat cctcctcctc ggagaagcag cgcttctact cctgcgaccg 6540
ctttgggggc cgtgagcccc cgaagcccaa gccctccctc agcagccacc caacgtcgcc 6600
aacagctggc caggagcggg gacccacccc acagggcagt ggttccgtga atgggagccc 6660
cttgctgtca acatctggtg ctagacccc cgcccgcggt gggcggaggc agctccccc 6720
gacgcccctg actccccgcc ccagcatcac ctacaagacg gccaaactct caccatcca 6780
cttcgccggg cttcagacca ccctccctgc ctttcccca ggccggctca gccgtgggct 6840
ttccgaacac aacgccctgc tgcagagaga cccctcagc cagccctgg cccctggctc 6900
tcgaattggc tctgaccctt acctggggca gcgtctggac agtgaggcct ctgtccacgc 6960
cctgcctgag gacacgtca ctttcgagga ggctgtggcc accaaactcg gccgtcctc 7020
caggacttcc tacgtgtcct ccctgacctc ccagtctcac cctctccgcc gcgtgcccaa 7080
cggttaccac tgcaccctgg gactcagctc ggttgggcca gcacggcaca gctaccacca 7140
ccctgaccaa gaccactggg gctagctgca ccgtgaccgc tcagacgcct gcattcgagc 7200
ggcgtgtgtt ccagtggatg agttttatca tccacacggg gcagtcggcc ctcgggggag 7260
gccttgccca ccttggtgag gctcctgtgg ccctccctc cctcttttac 7320
tctagacgac gaataaagcc ctgttgcttg agtgtacgta ccgc 7364

```

<210> 6
 <211> 2339
 <212> PRT
 <213> Homo Sapiens

```

<400> 6
Met Val Arg Phe Gly Asp Glu Leu Gly Gly Arg Tyr Gly Gly Pro Gly
1      5      10      15
Gly Gly Glu Arg Ala Arg Gly Gly Gly Ala Gly Gly Ala Gly Gly Pro
20      25      30
Gly Pro Gly Gly Leu Gln Pro Gly Gln Arg Val Leu Tyr Lys Gln Ser
35      40      45
Ile Ala Gln Arg Ala Arg Thr Met Ala Leu Tyr Asn Pro Ile Pro Val
50      55      60
Lys Gln Asn Cys Phe Thr Val Asn Arg Ser Leu Phe Val Phe Ser Glu
65      70      75      80
Asp Asn Val Val Arg Lys Tyr Ala Lys Arg Ile Thr Glu Trp Pro Pro
85      90      95
Phe Glu Tyr Met Ile Leu Ala Thr Ile Ile Ala Asn Cys Ile Val Leu
100     105     110
Ala Leu Glu Gln His Leu Pro Asp Gly Asp Lys Thr Pro Met Ser Glu
115     120     125
Arg Leu Asp Asp Thr Glu Pro Tyr Phe Ile Gly Ile Phe Cys Phe Glu
130     135     140
Ala Gly Ile Lys Ile Ile Ala Leu Gly Phe Val Phe His Lys Gly Ser
145     150     155     160
Tyr Leu Arg Asn Gly Trp Asn Val Met Asp Phe Val Val Val Leu Thr
165     170     175
Gly Ile Leu Ala Thr Ala Gly Thr Asp Phe Asp Leu Arg Thr Leu Arg
180     185     190
Ala Val Arg Val Leu Arg Pro Leu Lys Leu Val Ser Gly Ile Pro Ser
195     200     205

```

```

Leu Gln Val Val Leu Lys Ser Ile Met Lys Ala Met Val Pro Leu Leu
 210          215          220
Gln Ile Gly Leu Leu Leu Phe Phe Ala Ile Leu Met Phe Ala Ile Ile
 225          230          235          240
Gly Leu Glu Phe Tyr Met Gly Lys Phe His Lys Ala Cys Phe Pro Asn
          245          250          255
Ser Thr Asp Ala Glu Pro Val Gly Asp Phe Pro Cys Gly Lys Glu Ala
          260          265          270
Pro Ala Arg Leu Cys Glu Gly Asp Thr Glu Cys Arg Glu Tyr Trp Pro
          275          280          285
Gly Pro Asn Phe Gly Ile Thr Asn Phe Asp Asn Ile Leu Phe Ala Ile
          290          295          300
Leu Thr Val Phe Gln Cys Ile Thr Met Glu Gly Trp Thr Asp Ile Leu
 305          310          315          320
Tyr Asn Thr Asn Asp Ala Ala Gly Asn Thr Trp Asn Trp Leu Tyr Phe
          325          330          335
Ile Pro Leu Ile Ile Ile Gly Ser Phe Phe Met Leu Asn Leu Val Leu
          340          345          350
Gly Val Leu Ser Gly Glu Phe Ala Lys Glu Arg Glu Arg Val Glu Asn
          355          360          365
Arg Arg Ala Phe Leu Lys Leu Arg Arg Gln Gln Gln Ile Glu Arg Glu
 370          375          380
Leu Asn Gly Tyr Leu Glu Trp Ile Phe Lys Ala Glu Glu Val Met Leu
 385          390          395          400
Ala Glu Glu Asp Arg Asn Ala Glu Glu Lys Ser Pro Leu Asp Val Leu
          405          410          415
Lys Arg Ala Ala Thr Lys Lys Ser Arg Asn Asp Leu Ile His Ala Glu
          420          425          430
Glu Gly Glu Asp Arg Phe Ala Asp Leu Cys Ala Val Gly Ser Pro Phe
          435          440          445
Ala Arg Ala Ser Leu Lys Ser Gly Lys Thr Glu Ser Ser Ser Tyr Phe
          450          455          460
Arg Arg Lys Glu Lys Met Phe Arg Phe Phe Ile Arg Arg Met Val Lys
 465          470          475          480
Ala Gln Ser Phe Tyr Trp Val Val Leu Cys Val Val Ala Leu Asn Thr
          485          490          495
Leu Cys Val Ala Met Val His Tyr Asn Gln Pro Arg Arg Leu Thr Thr
          500          505          510
Thr Leu Tyr Phe Ala Glu Phe Val Phe Leu Gly Leu Phe Leu Thr Glu
          515          520          525
Met Ser Leu Lys Met Tyr Gly Leu Gly Pro Arg Ser Tyr Phe Arg Ser
          530          535          540
Ser Phe Asn Cys Phe Asp Phe Gly Val Ile Val Gly Ser Val Phe Glu
 545          550          555          560
Val Val Trp Ala Ala Ile Lys Pro Gly Ser Ser Phe Gly Ile Ser Val
          565          570          575
Leu Arg Ala Leu Arg Leu Leu Arg Ile Phe Lys Val Thr Lys Tyr Trp
          580          585          590
Ser Ser Leu Arg Asn Leu Val Val Ser Leu Leu Asn Ser Met Lys Ser
          595          600          605
Ile Ile Ser Leu Leu Phe Leu Leu Phe Leu Phe Ile Val Val Phe Ala
          610          615          620
Leu Leu Gly Met Gln Leu Phe Gly Gly Gln Phe Asn Phe Gln Asp Glu
 625          630          635          640
Thr Pro Thr Thr Asn Phe Asp Thr Phe Pro Ala Ala Ile Leu Thr Val
          645          650          655
Phe Gln Ile Leu Thr Gly Glu Asp Trp Asn Ala Val Met Tyr His Gly
          660          665          670
Ile Glu Ser Gln Gly Gly Val Ser Lys Gly Met Phe Ser Ser Phe Tyr
          675          680          685
Phe Ile Val Leu Thr Leu Phe Gly Asn Tyr Thr Leu Leu Asn Val Phe
 690          695          700

```

```

Leu Ala Ile Ala Val Asp Asn Leu Ala Asn Ala Gln Glu Leu Thr Lys
705          710          715          720
Asp Glu Glu Glu Met Glu Glu Ala Ala Asn Gln Lys Leu Ala Leu Gln
          725          730          735
Lys Ala Lys Glu Val Ala Glu Val Ser Pro Met Ser Ala Ala Asn Ile
          740          745          750
Ser Ile Ala Ala Arg Gln Gln Asn Ser Ala Lys Ala Arg Ser Val Trp
          755          760          765
Glu Gln Arg Ala Ser Gln Leu Arg Leu Gln Asn Leu Arg Ala Ser Cys
          770          775          780
Glu Ala Leu Tyr Ser Glu Met Asp Pro Glu Glu Arg Leu Arg Phe Ala
785          790          795          800
Thr Thr Arg His Leu Arg Pro Asp Met Lys Thr His Leu Asp Arg Pro
          805          810          815
Leu Val Val Glu Leu Gly Arg Asp Gly Ala Arg Gly Pro Val Gly Gly
          820          825          830
Lys Ala Arg Pro Glu Ala Ala Glu Ala Pro Glu Gly Val Asp Pro Pro
          835          840          845
Arg Arg His His Arg His Arg Asp Lys Asp Lys Thr Pro Ala Ala Gly
          850          855          860
Asp Gln Asp Arg Ala Glu Ala Pro Lys Ala Glu Ser Gly Glu Pro Gly
865          870          875          880
Ala Arg Glu Glu Arg Pro Arg Pro His Arg Ser His Ser Lys Glu Ala
          885          890          895
Ala Gly Pro Pro Glu Ala Arg Ser Glu Arg Gly Arg Gly Pro Gly Pro
          900          905          910
Glu Gly Gly Arg Arg His His Arg Arg Gly Ser Pro Glu Glu Ala Ala
          915          920          925
Glu Arg Glu Pro Arg Arg His Arg Ala His Arg His Gln Asp Pro Ser
          930          935          940
Lys Glu Cys Ala Gly Ala Lys Gly Glu Arg Arg Ala Arg His Arg Gly
945          950          955          960
Gly Pro Arg Ala Gly Pro Arg Glu Ala Glu Ser Gly Glu Glu Pro Ala
          965          970          975
Arg Arg His Arg Ala Arg His Lys Ala Gln Pro Ala His Glu Ala Val
          980          985          990
Glu Lys Glu Thr Thr Glu Lys Glu Ala Thr Glu Lys Glu Ala Glu Ile
          995          1000          1005
Val Glu Ala Asp Lys Glu Lys Glu Leu Arg Asn His Gln Pro Arg Glu
          1010          1015          1020
Pro His Cys Asp Leu Glu Thr Ser Gly Thr Val Thr Val Gly Pro Met
1025          1030          1035          1040
His Thr Leu Pro Ser Thr Cys Leu Gln Lys Val Glu Glu Gln Pro Glu
          1045          1050          1055
Asp Ala Asp Asn Gln Arg Asn Val Thr Arg Met Gly Ser Gln Pro Pro
          1060          1065          1070
Asp Pro Asn Thr Ile Val His Ile Pro Val Met Leu Thr Gly Pro Leu
          1075          1080          1085
Gly Glu Ala Thr Val Val Pro Ser Gly Asn Val Asp Leu Glu Ser Gln
          1090          1095          1100
Ala Glu Gly Lys Lys Glu Val Glu Ala Asp Asp Val Met Arg Ser Gly
1105          1110          1115          1120
Pro Arg Pro Ile Val Pro Tyr Ser Ser Met Phe Cys Leu Ser Pro Thr
          1125          1130          1135
Asn Leu Leu Arg Arg Phe Cys His Tyr Ile Val Thr Met Arg Tyr Phe
          1140          1145          1150
Glu Val Val Ile Leu Val Val Ile Ala Leu Ser Ser Ile Ala Leu Ala
          1155          1160          1165
Ala Glu Asp Pro Val Arg Thr Asp Ser Pro Arg Asn Asn Ala Leu Lys
          1170          1175          1180
Tyr Leu Asp Tyr Ile Phe Thr Gly Val Phe Thr Phe Glu Met Val Ile
1185          1190          1195          1200

```


Lys Met Ile Asp Leu Gly Leu Leu Leu His Pro Gly Ala Tyr Phe Arg
 1205 1210 1215
 Asp Leu Trp Asn Ile Leu Asp Phe Ile Val Val Ser Gly Ala Leu Val
 1220 1225 1230
 Ala Phe Ala Phe Ser Gly Ser Lys Gly Lys Asp Ile Asn Thr Ile Lys
 1235 1240 1245
 Ser Leu Arg Val Leu Arg Val Leu Arg Pro Leu Lys Thr Ile Lys Arg
 1250 1255 1260
 Leu Pro Lys Leu Lys Ala Val Phe Asp Cys Val Val Asn Ser Leu Lys
 1265 1270 1275 1280
 Asn Val Leu Asn Ile Leu Ile Val Tyr Met Leu Phe Met Phe Ile Phe
 1285 1290 1295
 Ala Val Ile Ala Val Gln Leu Phe Lys Gly Lys Phe Phe Tyr Cys Thr
 1300 1305 1310
 Asp Glu Ser Lys Glu Leu Glu Arg Asp Cys Arg Gly Gln Tyr Leu Asp
 1315 1320 1325
 Tyr Glu Lys Glu Glu Val Glu Ala Gln Pro Arg Gln Trp Lys Lys Tyr
 1330 1335 1340
 Asp Phe His Tyr Asp Asn Val Leu Trp Ala Leu Leu Thr Leu Phe Thr
 1345 1350 1355 1360
 Val Ser Thr Gly Glu Gly Trp Pro Met Val Leu Lys His Ser Val Asp
 1365 1370 1375
 Ala Thr Tyr Glu Glu Gln Gly Pro Ser Pro Gly Tyr Arg Met Glu Leu
 1380 1385 1390
 Ser Ile Phe Tyr Val Val Tyr Phe Val Val Phe Pro Phe Phe Val
 1395 1400 1405
 Asn Ile Phe Val Ala Leu Ile Ile Ile Thr Phe Gln Glu Gln Gly Asp
 1410 1415 1420
 Lys Val Met Ser Glu Cys Ser Leu Glu Lys Asn Glu Arg Ala Cys Ile
 1425 1430 1435 1440
 Asp Phe Ala Ile Ser Ala Lys Pro Leu Thr Arg Tyr Met Pro Gln Asn
 1445 1450 1455
 Arg Gln Ser Phe Gln Tyr Lys Thr Trp Thr Phe Val Val Ser Pro Pro
 1460 1465 1470
 Phe Glu Tyr Phe Ile Met Ala Met Ile Ala Leu Asn Thr Val Val Leu
 1475 1480 1485
 Met Met Lys Phe Tyr Asp Ala Pro Tyr Glu Tyr Glu Leu Met Leu Lys
 1490 1495 1500
 Cys Leu Asn Ile Val Phe Thr Ser Met Phe Ser Met Glu Cys Val Leu
 1505 1510 1515 1520
 Lys Ile Ile Ala Phe Gly Val Leu Asn Tyr Phe Arg Asp Ala Trp Asn
 1525 1530 1535
 Val Phe Asp Phe Val Thr Val Leu Gly Ser Ile Thr Asp Ile Leu Val
 1540 1545 1550
 Thr Glu Ile Ala Glu Thr Asn Asn Phe Ile Asn Leu Ser Phe Leu Arg
 1555 1560 1565
 Leu Phe Arg Ala Ala Arg Leu Ile Lys Leu Leu Arg Gln Gly Tyr Thr
 1570 1575 1580
 Ile Arg Ile Leu Leu Trp Thr Phe Val Gln Ser Phe Lys Ala Leu Pro
 1585 1590 1595 1600
 Tyr Val Cys Leu Leu Ile Ala Met Leu Phe Phe Ile Tyr Ala Ile Ile
 1605 1610 1615
 Gly Met Gln Val Phe Gly Asn Ile Ala Leu Asp Asp Asp Thr Ser Ile
 1620 1625 1630
 Asn Arg His Asn Asn Phe Arg Thr Phe Leu Gln Ala Leu Met Leu Leu
 1635 1640 1645
 Phe Arg Ser Ala Thr Gly Glu Ala Trp His Glu Ile Met Leu Ser Cys
 1650 1655 1660
 Leu Ser Asn Gln Ala Cys Asp Glu Gln Ala Asn Ala Thr Glu Cys Gly
 1665 1670 1675 1680
 Ser Asp Phe Ala Tyr Phe Tyr Phe Val Ser Phe Ile Phe Leu Cys Ser
 1685 1690 1695

Phe Leu Met Leu Asn Leu Phe Val Ala Val Ile Met Asp Asn Phe Glu
 1700 1705 1710
 Tyr Leu Thr Arg Asp Ser Ser Ile Leu Gly Pro His His Leu Asp Glu
 1715 1720 1725
 Phe Ile Arg Val Trp Ala Glu Tyr Asp Pro Ala Ala Cys Gly Arg Ile
 1730 1735 1740
 Ser Tyr Asn Asp Met Phe Glu Met Leu Lys His Met Ser Pro Pro Leu
 1745 1750 1755 1760
 Gly Leu Gly Lys Lys Cys Pro Ala Arg Val Ala Tyr Lys Arg Leu Val
 1765 1770 1775
 Arg Met Asn Met Pro Ile Ser Asn Glu Asp Met Thr Val His Phe Thr
 1780 1785 1790
 Ser Thr Leu Met Ala Leu Ile Arg Thr Ala Leu Glu Ile Lys Leu Ala
 1795 1800 1805
 Pro Ala Gly Thr Lys Gln His Gln Cys Asp Ala Glu Leu Arg Lys Glu
 1810 1815 1820
 Ile Ser Val Val Trp Ala Asn Leu Pro Gln Lys Thr Leu Asp Leu Leu
 1825 1830 1835 1840
 Val Pro Pro His Lys Pro Asp Glu Met Thr Val Gly Lys Val Tyr Ala
 1845 1850 1855
 Ala Leu Met Ile Phe Asp Phe Tyr Lys Gln Asn Lys Thr Thr Arg Asp
 1860 1865 1870
 Gln Met Gln Gln Ala Pro Gly Gly Leu Ser Gln Met Gly Pro Val Ser
 1875 1880 1885
 Leu Phe His Pro Leu Lys Ala Thr Leu Glu Gln Thr Gln Pro Ala Val
 1890 1895 1900
 Leu Arg Gly Ala Arg Val Phe Leu Arg Gln Lys Ser Ser Thr Ser Leu
 1905 1910 1915 1920
 Ser Asn Gly Gly Ala Ile Gln Asn Gln Glu Ser Gly Ile Lys Glu Ser
 1925 1930 1935
 Val Ser Trp Gly Thr Gln Arg Thr Gln Asp Ala Pro His Glu Ala Arg
 1940 1945 1950
 Pro Pro Leu Glu Arg Gly His Ser Thr Glu Ile Pro Val Gly Arg Ser
 1955 1960 1965
 Gly Ala Leu Ala Val Asp Val Gln Met Gln Ser Ile Thr Arg Arg Gly
 1970 1975 1980
 Pro Asp Gly Glu Pro Gln Pro Gly Leu Glu Ser Gln Gly Arg Ala Ala
 1985 1990 1995 2000
 Ser Met Pro Arg Leu Ala Ala Glu Thr Gln Pro Val Thr Asp Ala Ser
 2005 2010 2015
 Pro Met Lys Arg Ser Ile Ser Thr Leu Ala Gln Arg Pro Arg Gly Thr
 2020 2025 2030
 His Leu Cys Ser Thr Thr Pro Asp Arg Pro Pro Pro Ser Gln Ala Ser
 2035 2040 2045
 Ser His His His His His Arg Cys His Arg Arg Arg Asp Arg Lys Gln
 2050 2055 2060
 Arg Ser Leu Glu Lys Gly Pro Ser Leu Ser Ala Asp Met Asp Gly Ala
 2065 2070 2075 2080
 Pro Ser Ser Ala Val Gly Pro Gly Leu Pro Pro Gly Glu Gly Pro Thr
 2085 2090 2095
 Gly Cys Arg Arg Glu Arg Glu Arg Arg Gln Glu Arg Gly Arg Ser Gln
 2100 2105 2110
 Glu Arg Arg Gln Pro Ser Ser Ser Ser Ser Glu Lys Gln Arg Phe Tyr
 2115 2120 2125
 Ser Cys Asp Arg Phe Gly Gly Arg Glu Pro Pro Lys Pro Lys Pro Ser
 2130 2135 2140
 Leu Ser Ser His Pro Thr Ser Pro Thr Ala Gly Gln Glu Pro Gly Pro
 2145 2150 2155 2160
 His Pro Gln Gly Ser Gly Ser Val Asn Gly Ser Pro Leu Leu Ser Thr
 2165 2170 2175
 Ser Gly Ala Ser Thr Pro Gly Arg Gly Gly Arg Arg Gln Leu Pro Gln
 2180 2185 2190

Thr Pro Leu Thr Pro Arg Pro Ser Ile Thr Tyr Lys Thr Ala Asn Ser
 2195 2200 2205
 Ser Pro Ile His Phe Ala Gly Ala Gln Thr Ser Leu Pro Ala Phe Ser
 2210 2215 2220
 Pro Gly Arg Leu Ser Arg Gly Leu Ser Glu His Asn Ala Leu Leu Gln
 2225 2230 2235 2240
 Arg Asp Pro Leu Ser Gln Pro Leu Ala Pro Gly Ser Arg Ile Gly Ser
 2245 2250 2255
 Asp Pro Tyr Leu Gly Gln Arg Leu Asp Ser Glu Ala Ser Val His Ala
 2260 2265 2270
 Leu Pro Glu Asp Thr Leu Thr Phe Glu Glu Ala Val Ala Thr Asn Ser
 2275 2280 2285
 Gly Arg Ser Ser Arg Thr Ser Tyr Val Ser Ser Leu Thr Ser Gln Ser
 2290 2295 2300
 His Pro Leu Arg Arg Val Pro Asn Gly Tyr His Cys Thr Leu Gly Leu
 2305 2310 2315 2320
 Ser Ser Gly Gly Arg Ala Arg His Ser Tyr His His Pro Asp Gln Asp
 2325 2330 2335
 His Trp Cys

<210> 7
 <211> 7177
 <212> DNA
 <213> Homo Sapiens

<400> 7
 gcggcgccggg ctgcggcggtt ggggcccgggc gaggtccgct gcgggtcccgg cggctccgtg 60
 gctgctccgc tctgagcgcc tggcgcgccc cgcgccctcc ctgccggggc cgtggggccg 120
 gggatgcacg cggggcccgg gagccatggt ccgcttcggg gacgagctgg gcggccgcta 180
 tggaggcccc ggccggcgag agcgggcccgg gggcgccggg gccggcgggg cggggggccc 240
 ggggtcccggg gggctgcagc ccggccagcg ggtcctctac aagcaatcga tcgcgcagcg 300
 cgcgcggacc atggcgctgt acaaccccat ccgggtcaag cagaactgct tcaccgtcaa 360
 ccgctcgctc ttcgtcttca gcgaggacaa cgctcgccgc aaatacgcga agcgcatcac 420
 cgagtggcct ccattcgagt atatgatcct ggccaccatc atcgccaact gcatcggtgt 480
 ggccctggag cagcacctcc ctgatgggga caaacgccc atgtccgagc ggctggacga 540
 cacggagccc tatttcatcg ggatcttttg cttcgaggca gggatcaaaa tcatcgctct 600
 gggctttgtc ttccacaagg gctctttacct gcggaacggc tggaaactca tggacttcgt 660
 ggtcgctcac acaggatcc ttgccacggc ttcgacctgc gaacactgag 720
 ggctgtgcgt gtgctgaggg ccctgaagct ggtgtctggg attccaagtt tgcagggtgt 780
 gctcaagtcc atcatgaagg ccattgggtcc actcctgcag attgggctgc ttctcttctt 840
 tgccatcctc atgtttgcca tcattggcct ggagttctac atgggcaagt tccacaaggc 900
 ctgtttcccc aacagcacag atgcggagcg cgtgggtgac ttcccctgtg gcaaggaggc 960
 cccagcccgg ctgtgcgagg gcgacactga gtgccgggag tactggccag gacccaactt 1020
 tggcatcacc aactttgaca atatcctgtt tgccatcttg acggtgttcc agtgcacac 1080
 catggagggc tggactgaca tcctctataa tacaacgat gcggccggga acacctggaa 1140
 ctggctctac ttcateccct tcacatcat cggctccttc tcoatgctca acctgggtgt 1200
 gggcgtgctc tcgggggagt ttgccaaagg gcgagagagg gtggagaacc gccgcgcctt 1260
 cctgaagctg cgccggcagc agcagatcga gcgagagctc aacgggtacc tggagtggat 1320
 cttcaaggcg gaggaagtca tgctggccga ggaggacagg aatgcagagg agaagtcccc 1380
 tttggacgtg ctgaagagag cggccaccaa gaagagcaga aatgacctga tccacgcaga 1440
 ggaggggagag gaccggtttg cagatctctg tgctgttggg tcccccttcg cccgcgccag 1500
 cctcaagagc gggaagacag agagctcgct atacttccgg aggaaggaga agatgttccg 1560
 gttttttatc cggcgcatgg tgaaggctca gagcttctac tgggtgggtg tggtgcgtgt 1620
 ggccctgaac acactgtgtg tggccatggg gcatataaac cagccgcggc ggcttaccac 1680
 gaccctgtat tttgcagagt ttgttttctt cgggtctctt ctcacagaga tgtccctgaa 1740
 gatgtatggc ctggggccca gaagctactt ccggctcctc ttcaactgct tcgacttttg 1800
 ggtcatcgtg gggagcgtct ttgaagtggg ctggggcgcc atcaagccgg gaagctcctt 1860
 tgggcatcag gtgtgcggg ccctccgct gctgaggatc ttcaaagtca cgaagtact 1920
 gagctccctg cggaaacctg tgggtgtccct gctgaactcc atgaagtcca tcatcagcct 1980
 gctcttcttg ctcttcctgt tcattgttgt cttcgccctg ctgggggatg agctgttttg 2040
 gggacagttc aacttccagg atgagactcc cacaaccaac ttcgacacct tccctgccg 2100

catcctcact	gtcttccaga	tcctgacggg	agaggactgg	aatgcagtga	tgtatcacgg	2160
gatcgaatcg	caaggcgggc	tcagcaaagg	catgttctcg	tccttttact	tcattgtcct	2220
gacactgttc	ggaaactaca	ctctgctgaa	tgtctttctg	gccatcgctg	tggacaacct	2280
ggccaacgcc	caagagctga	ccaaggatga	agaggagatg	gaagaagcag	ccaatcagaa	2340
gcttgctctg	caaaaggcca	aagaagtggc	tgaagtcagc	cccatgtctg	ccgcgaacat	2400
ctccatcgcc	gccaggcgag	agaactcggc	caaggcgcg	tcggtgtggg	agcaggggc	2460
cagccagcta	cggctgcaga	acctgcgggc	cagctgcgag	gcgctgtaca	gcgagatgga	2520
ccccgaggag	cggctgcgct	tcgccactac	gcgccacctg	cggcccgaca	tgaagacgca	2580
cctggaccgg	ccgctggtgg	tggagctggg	ccgcgacggc	gcgcgggggc	ccgtgggagg	2640
caaagcccga	cccgaggctg	cggaggcccc	cgaggcgctc	gacctccgc	gcaggcacca	2700
ccggcaccgc	gacaaggaca	agacccccgc	ggcgggggac	caggaccgag	cagaggcccc	2760
gaaggcggag	agcggggagc	ccggtgcccc	ggaggagcgg	ccgcggccgc	accgcagcca	2820
cagcaaggag	gcccgggggc	ccccggaggc	gcggagcgag	cgcggccgag	gcccaggccc	2880
cgacccggag	cccgcgccac	accggcgcg	ctcccggag	gaggcgccg	agcggggacc	2940
ccgacgccac	cgcgcgcacc	ggcaccagga	tcgagcaag	gagtgcgcgc	gcgccaagg	3000
cgagcggcgc	gcgcggcacc	gcggcgcccc	ccgagcgggg	ccccgggagg	cggagagcgg	3060
ggaggagccg	gcgcggcgcc	accgggcccc	gcacaaggcg	cagcctgtct	acgaggctgt	3120
ggagaaggag	accacggaga	aggaggccac	ggagaaggag	gctgagatag	tggagaccga	3180
caaggaaaag	gagctccgga	accaccagcc	ccgggagcca	cactgtgacc	tggagaccag	3240
tgggactgtg	actgtgggtc	ccatgcacac	actgcccagc	acctgtctcc	agaagggtgga	3300
ggaacagcca	gaggatgcag	acaatcagcg	gaacgtcact	cgcatgggca	gtcagccccc	3360
agacccggag	atattgtac	atatccagc	gagctgacg	ggccctcttg	gggaagccac	3420
ggtcgttccc	agtggtaacg	tggacctgga	aagccaagca	gaggggaaga	aggagggtgga	3480
agcggatgac	gtgatgagga	gcggcccccg	gcctatcgct	ccatacagct	ccatgttctg	3540
tttaagcccc	accaacctgc	tcggccgctt	ctgccactac	atcgtgacca	tgaggtaact	3600
cgaggttgc	ttctcaggt	tcacgcctt	gagcgcatc	gccctggctg	ctgaggacc	3660
agtgcgcaca	gactcgcaca	ggaacaacgc	tctgaaatac	ctggattaca	tttccactgg	3720
tgtctttacc	tttgagatgg	tgataaagat	gatcgacttg	ggactgctgc	ttcaccctgg	3780
agcctatttc	cgggacttgt	ggaacattct	ggacttcat	gtggctcagt	gcgccctggt	3840
ggcgttctgc	ttctcaggt	ccaaaggga	agacatcaat	accatcaagt	ctctgagagt	3900
ccttcgtgtc	ctgcggcccc	tcaagaccat	caaacggctg	cccaagctca	aggctgtgtt	3960
tgactgtgtg	gtgaactccc	tgaagaatgt	cctcaacatc	ttgattgtct	acatgtctct	4020
catgttcata	tttgccgtca	ttgcgggtga	gctcttcaaa	gggaagtttt	tctactgcac	4080
agatgtatcc	aaggagctgg	agagggactg	caggggctag	tatttggatt	atgagaagga	4140
ggaagtggaa	gctcagccca	ggcagtgga	gaaatacgac	tttccactacg	acaatgtgct	4200
ctgggctctg	ctgacgctgt	tcacagtgct	cacgggagaa	ggctggccca	tgggtgctgaa	4260
acactccgtg	gatgccacct	atgaggagca	gggtccaaag	cctgggtacc	gcatggagct	4320
gtccatcttc	tacgtgtgtc	actttgtgtt	ctttcccttc	ttcttcgtca	acatctttgt	4380
ggctttgatc	atcatcacct	tcaggagca	gggggacaag	gtgatgtctg	aatgcagcct	4440
ggagaagaac	gagagggtct	gcattgactt	cgccatcagc	gccaaacccc	tgacacggta	4500
catgccccaa	aaccggcagt	cgttccagta	ttaagacgtg	acatttgtgg	tctccccgcc	4560
ctttgaatac	ttcatcatgg	ccatgatagc	cctcaacact	gtgggtgctga	tgatgaagtt	4620
ctatgatgca	ccctatgagt	acgagctgat	gctgaaatgc	ctgaacatcg	tgttcacatc	4680
catgttctcc	atggaatcgc	tgctgaagat	catcgctctt	gggggtgctga	actatttcag	4740
agatgcctgg	aatgtctttg	actttgtcac	tgtgttggga	agtattactg	atattttagt	4800
aacagagatt	gcggaacga	acaatttcat	caacctcagc	ttcctccgcc	tctttcgagc	4860
tgcgcggtcg	atcaagctgc	tcgcgccagg	ctacaccatc	cgcacccctg	tgtggacctt	4920
tgtccagtc	ttcaaggccc	tgccctacgt	gtgtctgctc	attgccatgc	tgttcttcat	4980
ctacgccatc	atcggcatgc	aggtgtttgg	gaatattgcc	ctggatgatg	acaccagcat	5040
caaccgccac	aacaacttcc	ggacgttttt	gcaagccctg	atgctgctgt	tcaggagcgc	5100
cacggggggag	gcctggcacg	agatcatgct	gtcctgcctg	agcaaccagg	cctgtgatga	5160
gcaggccaat	gccaccgagt	gtggaagtga	ctttgcctac	ttctacttcg	tctccttcat	5220
cttctgtg	tcctttctga	tgttgaacct	ctttgtggct	gtgatcatgg	acaattttga	5280
gtacctcagc	cgggactctt	ccatccctagg	tcctcaccac	ttggatgagt	tcacccgggt	5340
ctgggctgaa	tacgaccgg	ctgcgttgg	gcgcctcagt	tacaatgaca	tgtttgagat	5400
gctgaacac	atgtccccgc	ctctggggct	ggggaagaaa	tgccctgctc	gagttgctta	5460
caagcgcctg	gttcgcatga	acatgcccat	ctccaacgag	gacatgactg	ttcacttcac	5520
gtccacgctg	atggccctca	tccggacggc	actggagatc	aagctggccc	cagctgggac	5580
aaagcagcat	cagtgtgacg	cggagtggag	gaaggagatt	tccgttgtgt	gggccaatct	5640
gccccagaag	actttggact	tgctgggtacc	accccataag	cctgatgaga	tgacagtggg	5700
gaagggttat	gcagctctga	tgatatttga	cttctacaag	cagaacaaaa	ccaccagaga	5760
ccagatgcag	caggctcctg	gaggcctctc	ccagatgggt	cctgtgtccc	tgttccaccc	5820

tctgaaggcc	accctggagc	agacacagcc	ggctgtgctc	cgaggagccc	gggttttcct	5880
tcgacagaag	agttccacct	ccctcagcaa	tggcgggggc	atacaaaacc	aagagagtgg	5940
catcaaagag	tctgtctcct	ggggcactca	aaggaccctg	gatgcacccc	atgaggccag	6000
gccacccctg	gagcgtggcc	actccacaga	gatccctgtg	gggcggtcag	gagcactggc	6060
tgtggacgtt	cagatgcaga	gcataaccgc	gagggggcct	gatggggagc	cccagcctgg	6120
gctggagagc	cagggtcgag	cggcctccat	gccccgcctt	gcggccgaga	ctcagcccgt	6180
cacagatgcc	agcccatga	agcgctccat	ctccacgtcg	gcccagcggc	cccgtgggac	6240
tcattcttgc	agcaccaccc	cggaccgccc	accccctagc	caggcgtcgt	cgcaccacca	6300
ccaccaccgc	tgccaccgcc	gcagggacag	gaagcagagg	tccttgagga	agggggccag	6360
cctgtctgcc	gatatggatg	gcgcaccaag	cagtgtctgt	gggcccgggg	tgcccccggg	6420
agagggggcct	acaggctgcc	ggcgggaaag	agagcgccgg	caggagcggg	gccgggtcca	6480
ggagcggagg	cagccctcat	cctcctcctc	ggagaagcag	cgcttctact	cctgcgaccg	6540
ctttgggggg	cgtgagcccc	cgaagcccaa	gccctccctc	agcagccacc	caacgtcgcc	6600
aacagctggc	caggagccgg	gacccacccc	acaggccggc	tcagccgtgg	gctttccgaa	6660
cacaacgccc	tgctgcagag	agaccccttc	agccagcccc	tggcccctgg	ctctcgaatt	6720
ggctctgacc	cttacctggg	gcagcgtctg	gacagtggag	cctctgtcca	cgccctgcct	6780
gaggacacgc	tcacttttga	ggaggctgtg	gccaccaact	cgggccgctc	ctccaggact	6840
tcctacgtgt	cctccctgac	ctcccagctc	caccctctcc	gccgcgtgcc	caacggttac	6900
cactgcagccc	tgggactcag	ctcgggtggc	cgagcacggc	acagctacca	ccaccctgac	6960
caagaccact	ggtgctagct	gcaccgtgac	cgctcagacg	cctgcattga	gcaggcgtgt	7020
gttccagtgg	atgagtttta	tcattccacac	ggggcagtcg	gccctcgggg	gaggccttgc	7080
ccacettggt	gaggctcctg	tggccctccc	ctccccctcc	ccccctcttt	tactctagac	7140
gacgaataaa	gccctgttgc	ttgagtgtac	gtaccgc			7177

<210> 8
 <211> 2237
 <212> PRT
 <213> Homo Sapiens

<400> 8
 Met Val Arg Phe Gly Asp Glu Leu Gly Gly Arg Tyr Gly Gly Pro Gly
 1 5 10 15
 Gly Gly Glu Arg Ala Arg Gly Gly Ala Gly Ala Gly Gly Pro
 20 25 30
 Gly Pro Gly Gly Leu Gln Pro Gly Gln Arg Val Leu Tyr Lys Gln Ser
 35 40 45
 Ile Ala Gln Arg Ala Arg Thr Met Ala Leu Tyr Asn Pro Ile Pro Val
 50 55 60
 Lys Gln Asn Cys Phe Thr Val Asn Arg Ser Leu Phe Val Phe Ser Glu
 65 70 75 80
 Asp Asn Val Val Arg Lys Tyr Ala Lys Arg Ile Thr Glu Trp Pro Pro
 85 90 95
 Phe Glu Tyr Met Ile Leu Ala Thr Ile Ile Ala Asn Cys Ile Val Leu
 100 105 110
 Ala Leu Glu Gln His Leu Pro Asp Gly Asp Lys Thr Pro Met Ser Glu
 115 120 125
 Arg Leu Asp Asp Thr Glu Pro Tyr Phe Ile Gly Ile Phe Cys Phe Glu
 130 135 140
 Ala Gly Ile Lys Ile Ile Ala Leu Gly Phe Val Phe His Lys Gly Ser
 145 150 155 160
 Tyr Leu Arg Asn Gly Trp Asn Val Met Asp Phe Val Val Val Leu Thr
 165 170 175
 Gly Ile Leu Ala Thr Ala Gly Thr Asp Phe Asp Leu Arg Thr Leu Arg
 180 185 190
 Ala Val Arg Val Leu Arg Pro Leu Lys Leu Val Ser Gly Ile Pro Ser
 195 200 205
 Leu Gln Val Val Leu Lys Ser Ile Met Lys Ala Met Val Pro Leu Leu
 210 215 220
 Gln Ile Gly Leu Leu Leu Phe Phe Ala Ile Leu Met Phe Ala Ile Ile
 225 230 235 240

Gly	Leu	Glu	Phe	Tyr	Met	Gly	Lys	Phe	His	Lys	Ala	Cys	Phe	Pro	Asn
				245					250					255	
Ser	Thr	Asp	Ala	Glu	Pro	Val	Gly	Asp	Phe	Pro	Cys	Gly	Lys	Glu	Ala
			260					265					270		
Pro	Ala	Arg	Leu	Cys	Glu	Gly	Asp	Thr	Glu	Cys	Arg	Glu	Tyr	Trp	Pro
		275					280					285			
Gly	Pro	Asn	Phe	Gly	Ile	Thr	Asn	Phe	Asp	Asn	Ile	Leu	Phe	Ala	Ile
	290					295					300				
Leu	Thr	Val	Phe	Gln	Cys	Ile	Thr	Met	Glu	Gly	Trp	Thr	Asp	Ile	Leu
305					310					315					320
Tyr	Asn	Thr	Asn	Asp	Ala	Ala	Gly	Asn	Thr	Trp	Asn	Trp	Leu	Tyr	Phe
			325					330						335	
Ile	Pro	Leu	Ile	Ile	Ile	Gly	Ser	Phe	Met	Leu	Asn	Leu	Val	Leu	
			340					345					350		
Gly	Val	Leu	Ser	Gly	Glu	Phe	Ala	Lys	Glu	Arg	Glu	Arg	Val	Glu	Asn
		355					360					365			
Arg	Arg	Ala	Phe	Leu	Lys	Leu	Arg	Arg	Gln	Gln	Gln	Ile	Glu	Arg	Glu
	370					375					380				
Leu	Asn	Gly	Tyr	Leu	Glu	Trp	Ile	Phe	Lys	Ala	Glu	Glu	Val	Met	Leu
385					390					395					400
Ala	Glu	Glu	Asp	Arg	Asn	Ala	Glu	Glu	Lys	Ser	Pro	Leu	Asp	Val	Leu
			405						410					415	
Lys	Arg	Ala	Ala	Thr	Lys	Lys	Ser	Arg	Asn	Asp	Leu	Ile	His	Ala	Glu
			420					425					430		
Glu	Gly	Glu	Asp	Arg	Phe	Ala	Asp	Leu	Cys	Ala	Val	Gly	Ser	Pro	Phe
		435					440					445			
Ala	Arg	Ala	Ser	Leu	Lys	Ser	Gly	Lys	Thr	Glu	Ser	Ser	Ser	Tyr	Phe
	450					455					460				
Arg	Arg	Lys	Glu	Lys	Met	Phe	Arg	Phe	Phe	Ile	Arg	Arg	Met	Val	Lys
465					470					475					480
Ala	Gln	Ser	Phe	Tyr	Trp	Val	Val	Leu	Cys	Val	Val	Ala	Leu	Asn	Thr
			485					490						495	
Leu	Cys	Val	Ala	Met	Val	His	Tyr	Asn	Gln	Pro	Arg	Arg	Leu	Thr	Thr
			500					505					510		
Thr	Leu	Tyr	Phe	Ala	Glu	Phe	Val	Phe	Leu	Gly	Leu	Phe	Leu	Thr	Glu
		515					520					525			
Met	Ser	Leu	Lys	Met	Tyr	Gly	Leu	Gly	Pro	Arg	Ser	Tyr	Phe	Arg	Ser
	530					535					540				
Ser	Phe	Asn	Cys	Phe	Asp	Phe	Gly	Val	Ile	Val	Gly	Ser	Val	Phe	Glu
545					550					555					560
Val	Val	Trp	Ala	Ala	Ile	Lys	Pro	Gly	Ser	Ser	Phe	Gly	Ile	Ser	Val
			565					570						575	
Leu	Arg	Ala	Leu	Arg	Leu	Leu	Arg	Ile	Phe	Lys	Val	Thr	Lys	Tyr	Trp
			580					585					590		
Ser	Ser	Leu	Arg	Asn	Leu	Val	Val	Ser	Leu	Leu	Asn	Ser	Met	Lys	Ser
		595					600					605			
Ile	Ile	Ser	Leu	Leu	Phe	Leu	Phe	Leu	Phe	Ile	Val	Val	Phe	Ala	
	610					615				620					
Leu	Leu	Gly	Met	Gln	Leu	Phe	Gly	Gly	Gln	Phe	Asn	Phe	Gln	Asp	Glu
625					630					635					640
Thr	Pro	Thr	Thr	Asn	Phe	Asp	Thr	Phe	Pro	Ala	Ala	Ile	Leu	Thr	Val
			645						650					655	
Phe	Gln	Ile	Leu	Thr	Gly	Glu	Asp	Trp	Asn	Ala	Val	Met	Tyr	His	Gly
			660					665					670		
Ile	Glu	Ser	Gln	Gly	Gly	Val	Ser	Lys	Gly	Met	Phe	Ser	Ser	Phe	Tyr
		675					680					685			
Phe	Ile	Val	Leu	Thr	Leu	Phe	Gly	Asn	Tyr	Thr	Leu	Leu	Asn	Val	Phe
	690					695					700				
Leu	Ala	Ile	Ala	Val	Asp	Asn	Leu	Ala	Asn	Ala	Gln	Glu	Leu	Thr	Lys
705					710					715					720
Asp	Glu	Gly	Glu	Met	Glu	Glu	Ala	Ala	Asn	Gln	Lys	Leu	Ala	Leu	Gln
				725					730					735	

Lys Ala Lys Glu Val Ala Glu Val Ser Pro Met Ser Ala Ala Asn Ile
 740 745 750
 Ser Ile Ala Ala Arg Gln Gln Asn Ser Ala Lys Ala Arg Ser Val Trp
 755 760 765
 Glu Gln Arg Ala Ser Gln Leu Arg Leu Gln Asn Leu Arg Ala Ser Cys
 770 775 780
 Glu Ala Leu Tyr Ser Glu Met Asp Pro Glu Glu Arg Leu Arg Phe Ala
 785 790 795 800
 Thr Thr Arg His Leu Arg Pro Asp Met Lys Thr His Leu Asp Arg Pro
 805 810 815
 Leu Val Val Glu Leu Gly Arg Asp Gly Ala Arg Gly Pro Val Gly Gly
 820 825 830
 Lys Ala Arg Pro Glu Ala Ala Glu Ala Pro Glu Gly Val Asp Pro Pro
 835 840 845
 Arg Arg His His Arg His Arg Asp Lys Asp Lys Thr Pro Ala Ala Gly
 850 855 860
 Asp Gln Asp Arg Ala Glu Ala Pro Lys Ala Glu Ser Gly Glu Pro Gly
 865 870 875 880
 Ala Arg Glu Glu Arg Pro Arg Pro His Arg Ser His Ser Lys Glu Ala
 885 890 895
 Ala Gly Pro Pro Glu Ala Arg Ser Glu Arg Gly Arg Gly Pro Gly Pro
 900 905 910
 Glu Gly Gly Arg Arg His His Arg Arg Gly Ser Pro Glu Glu Ala Ala
 915 920 925
 Glu Arg Glu Pro Arg Arg His Arg Ala His Arg His Gln Asp Pro Ser
 930 935 940
 Lys Glu Cys Ala Gly Ala Lys Gly Glu Arg Arg Ala Arg His Arg Gly
 945 950 955 960
 Gly Pro Arg Ala Gly Pro Arg Glu Ala Glu Ser Gly Glu Glu Pro Ala
 965 970 975
 Arg Arg His Arg Ala Arg His Lys Ala Gln Pro Ala His Glu Ala Val
 980 985 990
 Glu Lys Glu Thr Thr Glu Lys Glu Ala Thr Glu Lys Glu Ala Glu Ile
 995 1000 1005
 Val Glu Ala Asp Lys Glu Lys Glu Leu Arg Asn His Gln Pro Arg Glu
 1010 1015 1020
 Pro His Cys Asp Leu Glu Thr Ser Gly Thr Val Thr Val Gly Pro Met
 1025 1030 1035 1040
 His Thr Leu Pro Ser Thr Cys Leu Gln Lys Val Glu Glu Gln Pro Glu
 1045 1050 1055
 Asp Ala Asp Asn Gln Arg Asn Val Thr Arg Met Gly Ser Gln Pro Pro
 1060 1065 1070
 Asp Pro Asn Thr Ile Val His Ile Pro Val Met Leu Thr Gly Pro Leu
 1075 1080 1085
 Gly Glu Ala Thr Val Val Pro Ser Gly Asn Val Asp Leu Glu Ser Gln
 1090 1095 1100
 Ala Glu Gly Lys Lys Glu Val Glu Ala Asp Asp Val Met Arg Ser Gly
 1105 1110 1115 1120
 Pro Arg Pro Ile Val Pro Tyr Ser Ser Met Phe Cys Leu Ser Pro Thr
 1125 1130 1135
 Asn Leu Leu Arg Arg Phe Cys His Tyr Ile Val Thr Met Arg Tyr Phe
 1140 1145 1150
 Glu Val Val Ile Leu Val Val Ile Ala Leu Ser Ser Ile Ala Leu Ala
 1155 1160 1165
 Ala Glu Asp Pro Val Arg Thr Asp Ser Pro Arg Asn Asn Ala Leu Lys
 1170 1175 1180
 Tyr Leu Asp Tyr Ile Phe Thr Gly Val Phe Thr Phe Glu Met Val Ile
 1185 1190 1195 1200
 Lys Met Ile Asp Leu Gly Leu Leu Leu His Pro Gly Ala Tyr Phe Arg
 1205 1210 1215
 Asp Leu Trp Asn Ile Leu Asp Phe Ile Val Val Ser Gly Ala Leu Val
 1220 1225 1230

Ala Phe Ala Phe Ser Gly Ser Lys Gly Lys Asp Ile Asn Thr Ile Lys
 1235 1240 1245
 Ser Leu Arg Val Leu Arg Val Leu Arg Pro Leu Lys Thr Ile Lys Arg
 1250 1255 1260
 Leu Pro Lys Leu Lys Ala Val Phe Asp Cys Val Val Asn Ser Leu Lys
 1265 1270 1275 1280
 Asn Val Leu Asn Ile Leu Ile Val Tyr Met Leu Phe Met Phe Ile Phe
 1285 1290 1295
 Ala Val Ile Ala Val Gln Leu Phe Lys Gly Lys Phe Phe Tyr Cys Thr
 1300 1305 1310
 Asp Glu Ser Lys Glu Leu Glu Arg Asp Cys Arg Gly Gln Tyr Leu Asp
 1315 1320 1325
 Tyr Glu Lys Glu Glu Val Glu Ala Gln Pro Arg Gln Trp Lys Lys Tyr
 1330 1335 1340
 Asp Phe His Tyr Asp Asn Val Leu Trp Ala Leu Leu Thr Leu Phe Thr
 1345 1350 1355 1360
 Val Ser Thr Gly Glu Gly Trp Pro Met Val Leu Lys His Ser Val Asp
 1365 1370 1375
 Ala Thr Tyr Glu Glu Gln Gly Pro Ser Pro Gly Tyr Arg Met Glu Leu
 1380 1385 1390
 Ser Ile Phe Tyr Val Val Tyr Phe Val Val Phe Pro Phe Phe Val
 1395 1400 1405
 Asn Ile Phe Val Ala Leu Ile Ile Ile Thr Phe Gln Glu Gln Gly Asp
 1410 1415 1420
 Lys Val Met Ser Glu Cys Ser Leu Glu Lys Asn Glu Arg Ala Cys Ile
 1425 1430 1435 1440
 Asp Phe Ala Ile Ser Ala Lys Pro Leu Thr Arg Tyr Met Pro Gln Asn
 1445 1450 1455
 Arg Gln Ser Phe Gln Tyr Lys Thr Trp Thr Phe Val Val Ser Pro Pro
 1460 1465 1470
 Phe Glu Tyr Phe Ile Met Ala Met Ile Ala Leu Asn Thr Val Val Leu
 1475 1480 1485
 Met Met Lys Phe Tyr Asp Ala Pro Tyr Glu Tyr Glu Leu Met Leu Lys
 1490 1495 1500
 Cys Leu Asn Ile Val Phe Thr Ser Met Phe Ser Met Glu Cys Val Leu
 1505 1510 1515 1520
 Lys Ile Ile Ala Phe Gly Val Leu Asn Tyr Phe Arg Asp Ala Trp Asn
 1525 1530 1535
 Val Phe Asp Phe Val Thr Val Leu Gly Ser Ile Thr Asp Ile Leu Val
 1540 1545 1550
 Thr Glu Ile Ala Glu Thr Asn Asn Phe Ile Asn Leu Ser Phe Leu Arg
 1555 1560 1565
 Leu Phe Arg Ala Ala Arg Leu Ile Lys Leu Leu Arg Gln Gly Tyr Thr
 1570 1575 1580
 Ile Arg Ile Leu Leu Trp Thr Phe Val Gln Ser Phe Lys Ala Leu Pro
 1585 1590 1595 1600
 Tyr Val Cys Leu Leu Ile Ala Met Leu Phe Phe Ile Tyr Ala Ile Ile
 1605 1610 1615
 Gly Met Gln Val Phe Gly Asn Ile Ala Leu Asp Asp Asp Thr Ser Ile
 1620 1625 1630
 Asn Arg His Asn Asn Phe Arg Thr Phe Leu Gln Ala Leu Met Leu Leu
 1635 1640 1645
 Phe Arg Ser Ala Thr Gly Glu Ala Trp His Glu Ile Met Leu Ser Cys
 1650 1655 1660
 Leu Ser Asn Gln Ala Cys Asp Glu Gln Ala Asn Ala Thr Glu Cys Gly
 1665 1670 1675 1680
 Ser Asp Phe Ala Tyr Phe Tyr Phe Val Ser Phe Ile Phe Leu Cys Ser
 1685 1690 1695
 Phe Leu Met Leu Asn Leu Phe Val Ala Val Ile Met Asp Asn Phe Glu
 1700 1705 1710
 Tyr Leu Thr Arg Asp Ser Ser Ile Leu Gly Pro His His Leu Asp Glu
 1715 1720 1725

Phe Ile Arg Val Trp Ala Glu Tyr Asp Pro Ala Ala Cys Gly Arg Ile
 1730 1735 1740
 Ser Tyr Asn Asp Met Phe Glu Met Leu Lys His Met Ser Pro Pro Leu
 1745 1750 1755 1760
 Gly Leu Gly Lys Lys Cys Pro Ala Arg Val Ala Tyr Lys Arg Leu Val
 1765 1770 1775
 Arg Met Asn Met Pro Ile Ser Asn Glu Asp Met Thr Val His Phe Thr
 1780 1785 1790
 Ser Thr Leu Met Ala Leu Ile Arg Thr Ala Leu Glu Ile Lys Leu Ala
 1795 1800 1805
 Pro Ala Gly Thr Lys Gln His Gln Cys Asp Ala Glu Leu Arg Lys Glu
 1810 1815 1820
 Ile Ser Val Val Trp Ala Asn Leu Pro Gln Lys Thr Leu Asp Leu Leu
 1825 1830 1835 1840
 Val Pro Pro His Lys Pro Asp Glu Met Thr Val Gly Lys Val Tyr Ala
 1845 1850 1855
 Ala Leu Met Ile Phe Asp Phe Tyr Lys Gln Asn Lys Thr Thr Arg Asp
 1860 1865 1870
 Gln Met Gln Gln Ala Pro Gly Gly Leu Ser Gln Met Gly Pro Val Ser
 1875 1880 1885
 Leu Phe His Pro Leu Lys Ala Thr Leu Glu Gln Thr Gln Pro Ala Val
 1890 1895 1900
 Leu Arg Gly Ala Arg Val Phe Leu Arg Gln Lys Ser Ser Thr Ser Leu
 1905 1910 1915 1920
 Ser Asn Gly Gly Ala Ile Gln Asn Gln Glu Ser Gly Ile Lys Glu Ser
 1925 1930 1935
 Val Ser Trp Gly Thr Gln Arg Thr Gln Asp Ala Pro His Glu Ala Arg
 1940 1945 1950
 Pro Pro Leu Glu Arg Gly His Ser Thr Glu Ile Pro Val Gly Arg Ser
 1955 1960 1965
 Gly Ala Leu Ala Val Asp Val Gln Met Gln Ser Ile Thr Arg Arg Gly
 1970 1975 1980
 Pro Asp Gly Glu Pro Gln Pro Gly Leu Glu Ser Gln Gly Arg Ala Ala
 1985 1990 1995 2000
 Ser Met Pro Arg Leu Ala Ala Glu Thr Gln Pro Val Thr Asp Ala Ser
 2005 2010 2015
 Pro Met Lys Arg Ser Ile Ser Thr Leu Ala Gln Arg Pro Arg Gly Thr
 2020 2025 2030
 His Leu Cys Ser Thr Thr Pro Asp Arg Pro Pro Pro Ser Gln Ala Ser
 2035 2040 2045
 Ser His His His His Arg Cys His Arg Arg Arg Asp Arg Lys Gln
 2050 2055 2060
 Arg Ser Leu Glu Lys Gly Pro Ser Leu Ser Ala Asp Met Asp Gly Ala
 2065 2070 2075 2080
 Pro Ser Ser Ala Val Gly Pro Gly Leu Pro Pro Gly Glu Gly Pro Thr
 2085 2090 2095
 Gly Cys Arg Arg Glu Arg Glu Arg Arg Gln Glu Arg Gly Arg Ser Gln
 2100 2105 2110
 Glu Arg Arg Gln Pro Ser Ser Ser Ser Ser Glu Lys Gln Arg Phe Tyr
 2115 2120 2125
 Ser Cys Asp Arg Phe Gly Gly Arg Glu Pro Pro Lys Pro Lys Pro Ser
 2130 2135 2140
 Leu Ser Ser His Pro Thr Ser Pro Thr Ala Gly Gln Glu Pro Gly Pro
 2145 2150 2155 2160
 His Pro Gln Ala Gly Ser Ala Val Gly Phe Pro Asn Thr Thr Pro Cys
 2165 2170 2175
 Cys Arg Glu Thr Pro Ser Ala Ser Pro Trp Pro Leu Ala Leu Glu Leu
 2180 2185 2190
 Ala Leu Thr Leu Thr Trp Gly Ser Val Trp Thr Val Arg Pro Leu Ser
 2195 2200 2205
 Thr Pro Cys Leu Arg Thr Arg Ser Leu Ser Arg Arg Leu Trp Pro Pro
 2210 2215 2220

Thr Arg Ala Ala Pro Pro Gly Leu Pro Thr Cys Pro Pro
2225 2230 2235

<210> 9
<211> 7808
<212> DNA
<213> Homo Sapiens

<400> 9
gatgtcccga gctgctatcc cgggctcggc cggggcagcc gccttctgag ccccgacc 60
gaggcgccga gccgcccgcg cccgatgggc tggggcgtgg agcgtctccg cagtcgtagc 120
tccagccgcc gcgctcccag ccccgccagc ctccagcatca gcggcgccgg cggcgccggc 180
ggcgtcttcc gcacgttccg ccgcagcgta acccgaggcc ctttctctct tgcagaatgg 240
cccgttccgg agacgagatg ccggcccgt acgggggagg aggtccggg gcagccggcg 300
gggtggctcg gggcagcgga ggccggcgag gagccgggg cagccggcag ggcgggcagc 360
ccggggcgca aaggatgtac aagcagtcga tggcgagag agcgccgacc atggcactct 420
acaaccccat ccccgctccga cagaactgcc tcacgggttaa ccggtctctc ttccctctca 480
gcgaagacaa cgtggtgaga aaatacgcca aaaagatcac cgaatggcct ccctttgaat 540
atatgatttt agccaccatc atagcgaatt gcactgctct cgcactggag cagcatctgc 600
ctgatgatga caagaccccg atgtctgaac ggttgatga cacagaacca tacttcattg 660
gaattttttg tttcgaggct ggaattaaaa tcattgccct tgggtttgcc ttccacaaag 720
gctcctactt gaggaatggc tggaaatgtca tggactttgt ggtggtgcta acgggcatct 780
tggcgacagt tgggacggag tttgacctac ggcagctgag ggcagttcga gtgctggggc 840
cgctcaagct ggtgtctgga atcccaagtt tacaagtcgt cctgaagtcg atcatgaagg 900
cgatgatccc tttgctgcag atcggcctcc tcctattttt tgcaatcctt atttttgcaa 960
tcatagggtt agaattttat atgggaaaaa ttcataccac ctgctttgaa gaggggacag 1020
atgacattca ggggtgagct cgggctccat gtgggacaga agagcccgcc cgcactgtcc 1080
ccaatgggac caaatgtcag ccctactggg aaggggccaa caacgggatc actcagttcg 1140
acaacatcct gtttgagtg ctgactgttt tcagtgcat aaccatggaa ggttgagactg 1200
atctcctact catcggtcc gatgcctcag ggaacacttg gaactggttg tacttcatcc 1260
ccctcatcat catcggtcc ttttttatgc tgaaccttgt gctgggtgtg ctgtcagggg 1320
agtttgccaa agaaagggaa cgggtggaga accggcgggc ttttctgaag ctgaggcgcc 1380
aacaacagat tgaacgtgag ctcaatgggt acatggaaat gatctcaaaa gcagaagagg 1440
tgatcctcgc cgaggtgaa actgacgggg agcagaggca tccctttgat ggagctctgc 1500
ggagaaccac cataaagaaa agcaagacag atttgctcaa cccgaagag gctgagatc 1560
agctgggtga tatagcctct gtgggttctc ccttcgccc agccagcatt aaaagtgcc 1620
agctggagaa ctgcaccttt tttcacaaaa aggagaggag gatgcgtttc tacatccgcc 1680
gcactgctca aactcaggcc ttctactgga ctgtactcag tttggtagct ctcaacacgc 1740
tgtgtgttgc tattgttcac tacaaccagc ccgagtggt ctccgacttc ctttactatg 1800
cagaattcat tttcttagga ctctttatgt ccgaaatgtt tataaaaaatg tacgggcttg 1860
ggacgcggcc ttacttccac tcttccctca actgctttga ctgtgggggt atcattggga 1920
gcatcttcga ggtcctcga gctgtcataa aaacctggca atcctttgga atcagcgtgt 1980
tacgagccct caggttattg cgtattttca aagtcacaa gtactgggca tctctcagaa 2040
acctggctgt ctctctcctc aactccatga agtccatcat cagcctgttg tttctccttt 2100
tcctgttcat tgtcgtcttc gcccttttgg gaatgcaact cttcgccggc cagtttaatt 2160
tcgatgaagg gactcctccc accaacttcg atacttttcc agcagcaata atgacggtgt 2220
ttcagatcct gacgggcaag gactggaaac aggtcatgta cgacgggatc aagtctcagg 2280
ggggcggtga gggcgccatg gtgttctcca tctatttcat tgtactgacg ctctttggga 2340
actacaccct cctgaatgtg ttcttggcca tcgctgtgga caatctggcc aacgcccagg 2400
agctcaccaa ggtggaggcg gacgagcaag aggaagaaga agcagcgaac cagaaacttg 2460
ccctacagaa agccaaggag gtggcagaag tgagtcctct gtccgcggcc aacatgtcta 2520
tagctgtgaa agagcaacag aagaatcaaa agccagccaa gtccgtgtgg gagcagcgga 2580
ccagtgaat gcgaaagcag aacttgctgg ccagccggga ggccctgtat aacgaaatgg 2640
acccggacga gcgctggaag gctgcctaca cggggcacct gggccagac atgaagacgc 2700
acttggaccg ccgctgggtg gtggaccgag aggagaccgc caacaacaac accaacaaga 2760
gcccggcgcc cgagcccacc gtggaccagc gcctcggcca gcagcgccgc gaggacttcc 2820
tcaggaaaca ggcccgtac cacgatcggg cccgggaccc cagcggtctg gcgggcttg 2880
acgcacggag gccctgggag ggaagccagg aggcgcagct gagccgggag ggaccctacg 2940
gcccgagtc gccaccacac gcccgggagg gcagcctgga gcaaccggg tctgggag 3000
gcgagggcca gcgaggcaag gccggggacc cccaccggag gcacgtgcac cggcagggg 3060
gcagcaggga gagccgcagc ggggtccccg gcacgggcgc ggacggggag catcgacgtc 3120
atcgcgcgca ccgcaggccc ggggaggagg gtccggagga caaggcgagg cggagggcg 3180

ggcaccgcga	gggcagccgg	ccggcccggg	gcggcgaggg	cgagggcgag	ggccccgacg	3240
ggggcgagcg	caggagaagg	caccggcatg	gcgctccagc	cacgtacgag	ggggacgcgc	3300
ggaggggagga	caaggagcgg	aggcatcggg	ggaggaaaga	gaaccagggc	tccggggtcc	3360
ctgtgtcggg	ccccaacctg	tcaaccaccc	ggccaatcca	gcaggacctg	ggccgccaag	3420
acccaccctt	ggcagaggat	attgacaaca	tgaagaacaa	caagctggcc	accgcggagt	3480
cgcccgctcc	ccacggcagc	cttgccacg	ccggcctgcc	ccagagccca	gccaaagatgg	3540
gaaacagcac	cgaccccgcc	cccatgctgg	ccatccctgc	catggccacc	aacccccaga	3600
acgccgccag	ccgccggacg	cccaacaacc	cggggaaccc	atccaatccc	ggccccccca	3660
agacccccga	gaatagcctt	atcgtaacca	accccagcgg	caccagacc	aattcagcta	3720
agactgccag	gaaacccgac	cacaccacag	tggacatccc	cccagcctgc	ccaccccccc	3780
tcaaccacac	cgctgtacaa	gtgaacaaaa	acgccaaccc	agaccactg	ccaaaaaaag	3840
aggaagagaa	gaaggaggag	gaggaagacg	accgtgggga	agacggccct	aagccaatgc	3900
ctccctatag	ctccatgttc	atcctgtcca	cgaccaaccc	ccttcgccgc	ctgtgccatt	3960
acatccgtga	ctcgcgtac	tttgagatgt	gcctcctcat	ggtcattgcc	atgagcgca	4020
tcgccctggc	cgccgaggac	cctgtgcagc	ccaacgcacc	tcggaacaac	gtgctgcgat	4080
actttgacta	cgtttttaca	ggcgtcttca	cctttgagat	ggtgatcaag	atgattgacc	4140
tggggctcgt	cctgcatcag	ggtgcctact	tccgtgacct	ctggaatatt	ctcgacttca	4200
tagtggtcag	tggggccctg	gtagcctttg	gtactcctgg	caatagcaaa	ggaaaagaca	4260
tcaacacgat	taaatccctc	cgagtccctc	gggtgctacg	acctcttaaa	accatcaagc	4320
ggctgccaaa	gctcaaggct	gtgtttgact	gtgtggtgaa	ctcacttaaa	aacgtcttca	4380
acatcctcat	cgtctacatg	ctattcatgt	tcatcttcgc	cgtggtggct	gtgcagctct	4440
tcaaggggaa	atctctccac	tgcactgacg	agtccaaaga	gtttgagaaa	gattgtcgag	4500
gcaaatacct	cctctacgag	aagaatgagg	tgaaggcgcg	agaccgggag	tggaaagaag	4560
atgaattcca	ttacgacaat	gtgctgtggg	ctctgctgac	cctcttcacc	gtgtccacgg	4620
gagaaggctg	gccacaggct	ctcaagcatt	cgggtggacgc	cacctttgag	aaccagggcc	4680
ccagccccgg	gtaccgcatg	gagatgtcca	ttttctacgt	cgtctacttt	gtgggtgttc	4740
ccatccctct	tgccaatatc	tttgtggcct	tgatcatcat	caccttccag	gagcaagggg	4800
acaagatgat	ggaggaatac	agcctggaga	aaaatgagag	ggcctgcatt	gatttcgcca	4860
tcagcgccaa	gccgctgacc	cgacacatgc	cgcagaacaa	gcagagcttc	cagtaccgca	4920
tgtggcagtt	cgtgggtgtc	ccgcctttcg	agtacacgat	catggccatg	atcgccctca	4980
acacatcgt	gcttatgatg	aagttctatg	tggtctctgt	tgcttatgaa	aatgccttgc	5040
gggtgttcaa	catcgtcttc	acctccctct	tctctctgga	atgtgtgctg	aaagtcatgg	5100
cttttgggat	tctgaattat	ttccgcgatg	cctggaacat	cttcgacttt	gtgactgttc	5160
tgggcagcat	caccgatatc	ctcgtgactg	agtttgggaa	tccgaataac	ttcatcaacc	5220
tgactcttct	ccgctcttcc	cgagctgccc	acttctccgt	acttctccgt	caggggtaca	5280
ccatccgcat	tcttctctgg	acctttgtgc	agtccttcaa	ggccctgcct	tatgtctgtc	5340
tgctgatcgc	catgctcttc	ttcatctatg	ccatcattgg	gatgcagggtg	tttggttaaca	5400
ttggcatcgc	cgtggaggac	gaggacagtg	atgaagatga	gttccaaatc	actgagcaca	5460
ataacttccg	gaccttcttc	caggccctca	tggtctctct	ccggagtgcc	accggggaag	5520
cttggcacia	catcatgctt	tcctgcctca	gcgggaaacc	gtgtgataag	aactctggca	5580
tcctgactcg	agagtgtggc	aatgaatttg	cttattttta	ctttgtttcc	ttcatcttcc	5640
tctgctcgtt	tctgatgctg	aatctctttg	tcgccgtcat	catggacaac	tttgagtacc	5700
tcacccgaga	ctctccatc	ctggggcccc	accacctgga	tgagtacgtg	cgtgtctggg	5760
ccgagtatga	ccccgcagct	tggggccgca	tgccctacct	ggacatgtat	cagatgctga	5820
gacacatgtc	tcggccctcg	ggtctgggga	agaagtgtcc	ggccagagtg	gcttacaagc	5880
ggcttctgcg	gatggacctg	cccgtcgcag	atgacaacac	cgtccacttc	aattccaccc	5940
tgcgcgagga	gatccgcaca	gccctggaca	tcaagattgc	caagggagga	gccgacaaac	6000
agcagatgga	cgctgagctg	cggaaggaga	tgatggcgat	ttggcccaat	ctgtcccaga	6060
agacgctaga	cctgctgggt	acacctcaca	agtcacagga	cctcacctg	gggaagatct	6120
acgcagccat	gatgatcatg	gagtactacc	ggcagagcaa	ggccaagaag	ctgcaggcca	6180
tgcgcgagga	gcaggaccgg	acacccctca	tgctccagcg	catggagccc	ccgtccccaa	6240
cgcaggaagg	gggacctggc	cagaacgccc	tcccctccac	ccagctggac	ccaggaggag	6300
ccctgatggc	tcacgaaagc	ggcctcaagg	agagcccgtc	ctgggtgacc	cagcgtgccc	6360
aggagatgtt	ccagaagacg	ggcacatgga	gtccggaaca	aggccccctt	accgacatgc	6420
ccaacagcca	gcctaactct	cagtcctggg	agatgcgaga	gatgggcaga	gatggctact	6480
ccgacagcga	gcaactacct	cccatggaag	gccagggccg	ggctgcctcc	atgccccgcc	6540
tccttgacga	gaaccagagg	agaaggggcc	ggccacgtgg	gaataacctc	agtaccatct	6600
cagacaccag	ccccatgaag	cgttcagcct	ccgtgctggg	ccccaaagcc	cgacgcctgg	6660
acgattactc	gctggagcgg	gtcccgcccg	agggaacca	gcggcaccac	cagcgggcgc	6720
gcgaccgcag	ccaccgcgcc	tctgagcgct	ccctgggccc	ctacaccgat	gtggacacag	6780
gcttggggac	agacctgagc	atgaccaccc	aatccgggga	cctgcctgct	aaggagcggg	6840
accaggagcg	gggcccggcc	aaggatcgga	agcatcgaca	gcaccaccac	caccaccacc	6900

```

accaccacca tccccgccc cccgacaagg accgctatgc ccaggaacgg ccggaccacg 6960
gccggggcacg ggctcgggac cagcgctggg cccgctcgcc cagcgagggc cgagagcaca 7020
tgggcgaccg gcagggcagt agttccgtaa gtggaagccc agccccctca acatctggta 7080
ccagcactcc gcggcggggc cgccgccagc tccccagac cccctccacc ccccgccac 7140
acgtgtccta ttcctctgtg atccgtaagg ccggcggtc ggggcccccg cagcagcagc 7200
agcagcagca gcagcagcag caggcggtgg ccaggccggg ccggcgggcc accagcggcc 7260
ctcggaggta cccaggcccc acggccgagc ctctggccgg agatcgggcg cccacggggg 7320
gccacagcag cggccgctcg cccaggatgg agaggcgggt cccaggcccc gcccgagcgg 7380
agtccccag ggctgtgcga cacggcgggg cccggtggcc ggcatctggc ccgcacgtgt 7440
ccgagggggc cccgggtccc cggcaccatg gctactacc gggctccgac tacgacgagg 7500
ccgatggccc gggcagcggg ggccgagagg aggccatggc cggggcctac gacgcgccac 7560
ccccgtacg acacgcgtcc tcgggcgcca ccggcgctc gccaggact cccggggcct 7620
cgggccccgg ctgcgcctcg ccttctcggc acggccggcg actccccaac ggctactacc 7680
cgggcgacgg atggccagg cccgcggggc cgggctccag gaagggcctg cagcaacct 7740
acagcgagag tgacgatgat tggtgctaag cccgggcgag gtggcgcccc cccggcccc 7800
cacgcacc

```

<210> 10
 <211> 2510
 <212> PRT
 <213> Homo Sapiens

```

<400> 10
Met Ala Arg Phe Gly Asp Glu Met Pro Ala Arg Tyr Gly Gly Gly Gly
1      5      10      15
Ser Gly Ala Ala Ala Gly Val Val Val Gly Ser Gly Gly Gly Arg Gly
20     25     30
Ala Gly Gly Ser Arg Gln Gly Gly Gln Pro Gly Ala Gln Arg Met Tyr
35     40     45
Lys Gln Ser Met Ala Gln Arg Ala Arg Thr Met Ala Leu Tyr Asn Pro
50     55     60
Ile Pro Val Arg Gln Asn Cys Leu Thr Val Asn Arg Ser Leu Phe Leu
65     70     75     80
Phe Ser Glu Asp Asn Val Val Arg Lys Tyr Ala Lys Lys Ile Thr Glu
85     90     95
Trp Pro Pro Phe Glu Tyr Met Ile Leu Ala Thr Ile Ile Ala Asn Cys
100    105    110
Ile Val Leu Ala Leu Glu Gln His Leu Pro Asp Asp Asp Lys Thr Pro
115    120    125
Met Ser Glu Arg Leu Asp Asp Thr Glu Pro Tyr Phe Ile Gly Ile Phe
130    135    140
Cys Phe Glu Ala Gly Ile Lys Ile Ile Ala Leu Gly Phe Ala Phe His
145    150    155    160
Lys Gly Ser Tyr Leu Arg Asn Gly Trp Asn Val Met Asp Phe Val Val
165    170    175
Val Leu Thr Gly Ile Leu Ala Thr Val Gly Thr Glu Phe Asp Leu Arg
180    185    190
Thr Leu Arg Ala Val Arg Val Leu Arg Pro Leu Lys Leu Val Ser Gly
195    200    205
Ile Pro Ser Leu Gln Val Val Leu Lys Ser Ile Met Lys Ala Met Ile
210    215    220
Pro Leu Leu Gln Ile Gly Leu Leu Leu Phe Phe Ala Ile Leu Ile Phe
225    230    235    240
Ala Ile Ile Gly Leu Glu Phe Tyr Met Gly Lys Phe His Thr Thr Cys
245    250    255
Phe Glu Glu Gly Thr Asp Asp Ile Gln Gly Glu Ser Pro Ala Pro Cys
260    265    270
Gly Thr Glu Glu Pro Ala Arg Thr Cys Pro Asn Gly Thr Lys Cys Gln
275    280    285
Pro Tyr Trp Glu Gly Pro Asn Asn Gly Ile Thr Gln Phe Asp Asn Ile

```

290	295	300
Leu Phe Ala Val	Leu Thr Val Phe Gln Cys Ile Thr Met Glu Gly Trp	
305	310	315
Thr Asp Leu Leu Tyr Asn Ser Asn Asp Ala Ser Gly Asn Thr Trp Asn		320
	325	330
Trp Leu Tyr Phe	Ile Pro Leu Ile Ile Ile Gly Ser Phe Phe Met Leu	335
	340	345
Asn Leu Val Leu Gly Val Leu Ser Gly Glu Phe Ala Lys Glu Arg Glu		350
	355	360
Arg Val Glu Asn Arg Arg Ala Phe Leu Lys Leu Arg Arg Gln Gln Gln		365
	370	375
Ile Glu Arg Glu Leu Asn Gly Tyr Met Glu Trp Ile Ser Lys Ala Glu		380
	385	390
Glu Val Ile Leu Ala Glu Asp Glu Thr Asp Gly Glu Gln Arg His Pro		395
	405	410
Phe Asp Gly Ala Leu Arg Arg Thr Thr Ile Lys Lys Ser Lys Thr Asp		415
	420	425
Leu Leu Asn Pro Glu Glu Ala Glu Asp Gln Leu Ala Asp Ile Ala Ser		430
	435	440
Val Gly Ser Pro Phe Ala Arg Ala Ser Ile Lys Ser Ala Lys Leu Glu		445
	450	455
Asn Ser Thr Phe Phe His Lys Lys Glu Arg Arg Met Arg Phe Tyr Ile		460
	465	470
Arg Arg Met Val Lys Thr Gln Ala Phe Tyr Trp Thr Val Leu Ser Leu		475
	485	490
Val Ala Leu Asn Thr Leu Cys Val Ala Ile Val His Tyr Asn Gln Pro		495
	500	505
Glu Trp Leu Ser Asp Phe Leu Tyr Tyr Ala Glu Phe Ile Phe Leu Gly		510
	515	520
Leu Phe Met Ser Glu Met Phe Ile Lys Met Tyr Gly Leu Gly Thr Arg		525
	530	535
Pro Tyr Phe His Ser Ser Phe Asn Cys Phe Asp Cys Gly Val Ile Ile		540
	545	550
Gly Ser Ile Phe Glu Val Ile Trp Ala Val Ile Lys Pro Gly Thr Ser		555
	565	570
Phe Gly Ile Ser Val Leu Arg Ala Leu Arg Leu Leu Arg Ile Phe Lys		575
	580	585
Val Thr Lys Tyr Trp Ala Ser Leu Arg Asn Leu Val Val Ser Leu Leu		590
	595	600
Asn Ser Met Lys Ser Ile Ile Ser Leu Leu Phe Leu Leu Phe Leu Phe		605
	610	615
Ile Val Val Phe Ala Leu Leu Gly Met Gln Leu Phe Gly Gly Gln Phe		620
	625	630
Asn Phe Asp Glu Gly Thr Pro Pro Thr Asn Phe Asp Thr Phe Pro Ala		635
	645	650
Ala Ile Met Thr Val Phe Gln Ile Leu Thr Gly Glu Asp Trp Asn Glu		655
	660	665
Val Met Tyr Asp Gly Ile Lys Ser Gln Gly Gly Val Gln Gly Gly Met		670
	675	680
Val Phe Ser Ile Tyr Phe Ile Val Leu Thr Leu Phe Gly Asn Tyr Thr		685
	690	695
Leu Leu Asn Val Phe Leu Ala Ile Ala Val Asp Asn Leu Ala Asn Ala		700
	705	710
Gln Glu Leu Thr Lys Val Glu Ala Asp Glu Gln Glu Glu Glu Ala		715
	725	730
Ala Asn Gln Lys Leu Ala Leu Gln Lys Ala Lys Glu Val Ala Glu Val		735
	740	745
Ser Pro Leu Ser Ala Ala Asn Met Ser Ile Ala Val Lys Glu Gln Gln		750
	755	760
Lys Asn Gln Lys Pro Ala Lys Ser Val Trp Glu Gln Arg Thr Ser Glu		765
	770	775
Met Arg Lys Gln Asn Leu Leu Ala Ser Arg Glu Ala Leu Tyr Asn Glu		780

```

785              790              795              800
Met Asp Pro Asp Glu Arg Trp Lys Ala Ala Tyr Thr Arg His Leu Arg
805              810              815
Pro Asp Met Lys Thr His Leu Asp Arg Pro Leu Val Val Asp Pro Gln
820              825              830
Glu Asn Arg Asn Asn Thr Asn Lys Ser Arg Ala Ala Glu Pro Thr
835              840              845
Val Asp Gln Arg Leu Gly Gln Gln Arg Ala Glu Asp Phe Leu Arg Lys
850              855              860
Gln Ala Arg Tyr His Asp Arg Ala Arg Asp Pro Ser Gly Ser Ala Gly
865              870              875
Leu Asp Ala Arg Arg Pro Trp Ala Gly Ser Gln Glu Ala Glu Leu Ser
885              890              895
Arg Glu Gly Pro Tyr Gly Arg Glu Ser Asp His His Ala Arg Glu Gly
900              905              910
Ser Leu Glu Gln Pro Gly Phe Trp Glu Gly Glu Ala Glu Arg Gly Lys
915              920              925
Ala Gly Asp Pro His Arg Arg His Val His Arg Gln Gly Gly Ser Arg
930              935              940
Glu Ser Arg Ser Gly Ser Pro Arg Thr Gly Ala Asp Gly Glu His Arg
945              950              955
Arg His Arg Ala His Arg Arg Pro Gly Glu Gly Pro Glu Asp Lys
965              970              975
Ala Glu Arg Arg Ala Arg His Arg Glu Gly Ser Arg Pro Ala Arg Gly
980              985              990
Gly Glu Gly Glu Gly Glu Gly Pro Asp Gly Gly Glu Arg Arg Arg Arg
995              1000              1005
His Arg His Gly Ala Pro Ala Thr Tyr Glu Gly Asp Ala Arg Arg Glu
1010              1015              1020
Asp Lys Glu Arg Arg His Arg Arg Arg Lys Glu Asn Gln Gly Ser Gly
1025              1030              1035
Val Pro Val Ser Gly Pro Asn Leu Ser Thr Thr Arg Pro Ile Gln Gln
1045              1050              1055
Asp Leu Gly Arg Gln Asp Pro Pro Leu Ala Glu Asp Ile Asp Asn Met
1060              1065              1070
Lys Asn Asn Lys Leu Ala Thr Ala Glu Ser Ala Ala Pro His Gly Ser
1075              1080              1085
Leu Gly His Ala Gly Leu Pro Gln Ser Pro Ala Lys Met Gly Asn Ser
1090              1095              1100
Thr Asp Pro Gly Pro Met Leu Ala Ile Pro Ala Met Ala Thr Asn Pro
1105              1110              1115
Gln Asn Ala Ala Ser Arg Arg Thr Pro Asn Asn Pro Gly Asn Pro Ser
1125              1130              1135
Asn Pro Gly Pro Pro Lys Thr Pro Glu Asn Ser Leu Ile Val Thr Asn
1140              1145              1150
Pro Ser Gly Thr Gln Thr Asn Ser Ala Lys Thr Ala Arg Lys Pro Asp
1155              1160              1165
His Thr Thr Val Asp Ile Pro Pro Ala Cys Pro Pro Pro Leu Asn His
1170              1175              1180
Thr Val Val Gln Val Asn Lys Asn Ala Asn Pro Asp Pro Leu Pro Lys
1185              1190              1195
Lys Glu Glu Glu Lys Lys Glu Glu Glu Glu Asp Asp Arg Gly Glu Asp
1205              1210              1215
Gly Pro Lys Pro Met Pro Pro Tyr Ser Ser Met Phe Ile Leu Ser Thr
1220              1225              1230
Thr Asn Pro Leu Arg Arg Leu Cys His Tyr Ile Leu Asn Leu Arg Tyr
1235              1240              1245
Phe Glu Met Cys Ile Leu Met Val Ile Ala Met Ser Ser Ile Ala Leu
1250              1255              1260
Ala Ala Glu Asp Pro Val Gln Pro Asn Ala Pro Arg Asn Asn Val Leu
1265              1270              1275
Arg Tyr Phe Asp Tyr Val Phe Thr Gly Val Phe Thr Phe Glu Met Val

```

			1285				1290				1295				
Ile	Lys	Met	Ile	Asp	Leu	Gly	Leu	Val	Leu	His	Gln	Gly	Ala	Tyr	Phe
			1300				1305						1310		
Arg	Asp	Leu	Trp	Asn	Ile	Leu	Asp	Phe	Ile	Val	Val	Ser	Gly	Ala	Leu
			1315				1320						1325		
Val	Ala	Phe	Ala	Phe	Thr	Gly	Asn	Ser	Lys	Gly	Lys	Asp	Ile	Asn	Thr
			1330				1335						1340		
Ile	Lys	Ser	Leu	Arg	Val	Leu	Arg	Val	Leu	Arg	Pro	Leu	Lys	Thr	Ile
			1345				1350						1355		1360
Lys	Arg	Leu	Pro	Lys	Leu	Lys	Ala	Val	Phe	Asp	Cys	Val	Val	Asn	Ser
			1365							1370					1375
Leu	Lys	Asn	Val	Phe	Asn	Ile	Leu	Ile	Val	Tyr	Met	Leu	Phe	Met	Phe
			1380							1385					1390
Ile	Phe	Ala	Val	Val	Ala	Val	Gln	Leu	Phe	Lys	Gly	Lys	Phe	Phe	His
			1395				1400						1405		
Cys	Thr	Asp	Glu	Ser	Lys	Glu	Phe	Glu	Lys	Asp	Cys	Arg	Gly	Lys	Tyr
			1410				1415						1420		
Leu	Leu	Tyr	Glu	Lys	Asn	Glu	Val	Lys	Ala	Arg	Asp	Arg	Glu	Trp	Lys
			1425				1430						1435		1440
Lys	Tyr	Glu	Phe	His	Tyr	Asp	Asn	Val	Leu	Trp	Ala	Leu	Leu	Thr	Leu
			1445							1450					1455
Phe	Thr	Val	Ser	Thr	Gly	Glu	Gly	Trp	Pro	Gln	Val	Leu	Lys	His	Ser
			1460							1465					1470
Val	Asp	Ala	Thr	Phe	Glu	Asn	Gln	Gly	Pro	Ser	Pro	Gly	Tyr	Arg	Met
			1475							1480					1485
Glu	Met	Ser	Ile	Phe	Tyr	Val	Val	Tyr	Phe	Val	Val	Phe	Pro	Phe	Phe
			1490				1495						1500		
Phe	Val	Asn	Ile	Phe	Val	Ala	Leu	Ile	Ile	Ile	Thr	Phe	Gln	Glu	Gln
			1505				1510						1515		1520
Gly	Asp	Lys	Met	Met	Glu	Glu	Tyr	Ser	Leu	Glu	Lys	Asn	Glu	Arg	Ala
			1525							1530					1535
Cys	Ile	Asp	Phe	Ala	Ile	Ser	Ala	Lys	Pro	Leu	Thr	Arg	His	Met	Pro
			1540							1545					1550
Gln	Asn	Lys	Gln	Ser	Phe	Gln	Tyr	Arg	Met	Trp	Gln	Phe	Val	Val	Ser
			1555				1560						1565		
Pro	Pro	Phe	Glu	Tyr	Thr	Ile	Met	Ala	Met	Ile	Ala	Leu	Asn	Thr	Ile
			1570				1575						1580		
Val	Leu	Met	Met	Lys	Phe	Tyr	Gly	Ala	Ser	Val	Ala	Tyr	Glu	Asn	Ala
			1585				1590						1595		1600
Leu	Arg	Val	Phe	Asn	Ile	Val	Phe	Thr	Ser	Leu	Phe	Ser	Leu	Glu	Cys
			1605							1610					1615
Val	Leu	Lys	Val	Met	Ala	Phe	Gly	Ile	Leu	Asn	Tyr	Phe	Arg	Asp	Ala
			1620							1625					1630
Trp	Asn	Ile	Phe	Asp	Phe	Val	Thr	Val	Leu	Gly	Ser	Ile	Thr	Asp	Ile
			1635							1640					1645
Leu	Val	Thr	Glu	Phe	Gly	Asn	Pro	Asn	Asn	Phe	Ile	Asn	Leu	Ser	Phe
			1650				1655						1660		
Leu	Arg	Leu	Phe	Arg	Ala	Ala	Arg	Leu	Ile	Lys	Leu	Leu	Arg	Gln	Gly
			1665				1670						1675		1680
Tyr	Thr	Ile	Arg	Ile	Leu	Leu	Trp	Thr	Phe	Val	Gln	Ser	Phe	Lys	Ala
			1685							1690					1695
Leu	Pro	Tyr	Val	Cys	Leu	Leu	Ile	Ala	Met	Leu	Phe	Phe	Ile	Tyr	Ala
			1700							1705					1710
Ile	Ile	Gly	Met	Gln	Val	Phe	Gly	Asn	Ile	Gly	Ile	Asp	Val	Glu	Asp
			1715							1720					1725
Glu	Asp	Ser	Asp	Glu	Asp	Glu	Phe	Gln	Ile	Thr	Glu	His	Asn	Asn	Phe
			1730				1735						1740		
Arg	Thr	Phe	Phe	Gln	Ala	Leu	Met	Leu	Leu	Phe	Arg	Ser	Ala	Thr	Gly
			1745				1750						1755		1760
Glu	Ala	Trp	His	Asn	Ile	Met	Leu	Ser	Cys	Leu	Ser	Gly	Lys	Pro	Cys
			1765							1770					1775
Asp	Lys	Asn	Ser	Gly	Ile	Leu	Thr	Arg	Glu	Cys	Gly	Asn	Glu	Phe	Ala

			1780					1785				1790			
Tyr	Phe	Tyr	Phe	Val	Ser	Phe	Ile	Phe	Leu	Cys	Ser	Phe	Leu	Met	Leu
		1795					1800					1805			
Asn	Leu	Phe	Val	Ala	Val	Ile	Met	Asp	Asn	Phe	Glu	Tyr	Leu	Thr	Arg
	1810					1815					1820				
Asp	Ser	Ser	Ile	Leu	Gly	Pro	His	His	Leu	Asp	Glu	Tyr	Val	Arg	Val
1825					1830					1835					1840
Trp	Ala	Glu	Tyr	Asp	Pro	Ala	Ala	Trp	Gly	Arg	Met	Pro	Tyr	Leu	Asp
				1845					1850					1855	
Met	Tyr	Gln	Met	Leu	Arg	His	Met	Ser	Pro	Pro	Leu	Gly	Leu	Gly	Lys
			1860					1865				1870			
Lys	Cys	Pro	Ala	Arg	Val	Ala	Tyr	Lys	Arg	Leu	Leu	Arg	Met	Asp	Leu
		1875					1880					1885			
Pro	Val	Ala	Asp	Asp	Asn	Thr	Val	His	Phe	Asn	Ser	Thr	Leu	Met	Ala
	1890					1895					1900				
Leu	Ile	Arg	Thr	Ala	Leu	Asp	Ile	Lys	Ile	Ala	Lys	Gly	Gly	Ala	Asp
1905					1910					1915					1920
Lys	Gln	Gln	Met	Asp	Ala	Glu	Leu	Arg	Lys	Glu	Met	Met	Ala	Ile	Trp
				1925					1930					1935	
Pro	Asn	Leu	Ser	Gln	Lys	Thr	Leu	Asp	Leu	Leu	Val	Thr	Pro	His	Lys
		1940						1945				1950			
Ser	Thr	Asp	Leu	Thr	Val	Gly	Lys	Ile	Tyr	Ala	Ala	Met	Met	Ile	Met
		1955					1960					1965			
Glu	Tyr	Tyr	Arg	Gln	Ser	Lys	Ala	Lys	Lys	Leu	Gln	Ala	Met	Arg	Glu
	1970					1975				1980					
Glu	Gln	Asp	Arg	Thr	Pro	Leu	Met	Phe	Gln	Arg	Met	Glu	Pro	Pro	Ser
1985					1990					1995					2000
Pro	Thr	Gln	Glu	Gly	Gly	Pro	Gly	Gln	Asn	Ala	Leu	Pro	Ser	Thr	Gln
				2005				2010						2015	
Leu	Asp	Pro	Gly	Gly	Ala	Leu	Met	Ala	His	Glu	Ser	Gly	Leu	Lys	Glu
			2020					2025					2030		
Ser	Pro	Ser	Trp	Val	Thr	Gln	Arg	Ala	Gln	Glu	Met	Phe	Gln	Lys	Thr
		2035					2040					2045			
Gly	Thr	Trp	Ser	Pro	Glu	Gln	Gly	Pro	Pro	Thr	Asp	Met	Pro	Asn	Ser
	2050					2055					2060				
Gln	Pro	Asn	Ser	Gln	Ser	Val	Glu	Met	Arg	Glu	Met	Gly	Arg	Asp	Gly
2065					2070					2075					2080
Tyr	Ser	Asp	Ser	Glu	His	Tyr	Leu	Pro	Met	Glu	Gly	Gln	Gly	Arg	Ala
				2085					2090					2095	
Ala	Ser	Met	Pro	Arg	Leu	Pro	Ala	Glu	Asn	Gln	Arg	Arg	Arg	Gly	Arg
			2100					2105				2110			
Pro	Arg	Gly	Asn	Asn	Leu	Ser	Thr	Ile	Ser	Asp	Thr	Ser	Pro	Met	Lys
		2115					2120					2125			
Arg	Ser	Ala	Ser	Val	Leu	Gly	Pro	Lys	Ala	Arg	Arg	Leu	Asp	Asp	Tyr
		2130				2135					2140				
Ser	Leu	Glu	Arg	Val	Pro	Glu	Glu	Asn	Gln	Arg	His	His	Gln	Arg	
2145					2150				2155					2160	
Arg	Arg	Asp	Arg	Ser	His	Arg	Ala	Ser	Glu	Arg	Ser	Leu	Gly	Arg	Tyr
			2165						2170					2175	
Thr	Asp	Val	Asp	Thr	Gly	Leu	Gly	Thr	Asp	Leu	Ser	Met	Thr	Thr	Gln
		2180						2185					2190		
Ser	Gly	Asp	Leu	Pro	Ser	Lys	Glu	Arg	Asp	Gln	Glu	Arg	Gly	Arg	Pro
		2195					2200					2205			
Lys	Asp	Arg	Lys	His	Arg	Gln	His	His	His	His	His	His	His	His	His
	2210					2215						2220			
His	Pro	Pro	Pro	Pro	Asp	Lys	Asp	Arg	Tyr	Ala	Gln	Glu	Arg	Pro	Asp
2225					2230					2235					2240
His	Gly	Arg	Ala	Arg	Ala	Arg	Asp	Gln	Arg	Trp	Ser	Arg	Ser	Pro	Ser
			2245						2250					2255	
Glu	Gly	Arg	Glu	His	Met	Ala	His	Arg	Gln	Gly	Ser	Ser	Ser	Val	Ser
			2260					2265					2270		
Gly	Ser	Pro	Ala	Pro	Ser	Thr	Ser	Gly	Thr	Ser	Thr	Pro	Arg	Arg	Gly

2275	2280	2285
Arg Arg Gln Leu Pro Gln Thr Pro Ser Thr Pro Arg Pro His Val Ser		
2290	2295	2300
Tyr Ser Pro Val Ile Arg Lys Ala Gly Gly Ser Gly Pro Pro Gln Gln		
2305	2310	2315
Gln Gln Gln Gln Gln Gln Gln Gln Ala Val Ala Arg Pro Gly Arg		2320
	2325	2330
Ala Ala Thr Ser Gly Pro Arg Arg Tyr Pro Gly Pro Thr Ala Glu Pro		2335
2340	2345	2350
Leu Ala Gly Asp Arg Pro Pro Thr Gly Gly His Ser Ser Gly Arg Ser		2365
2355	2360	2365
Pro Arg Met Glu Arg Arg Val Pro Gly Pro Ala Arg Ser Glu Ser Pro		2380
2370	2375	2380
Arg Ala Cys Arg His Gly Gly Ala Arg Trp Pro Ala Ser Gly Pro His		2400
2385	2390	2395
Val Ser Glu Gly Pro Pro Gly Pro Arg His His Gly Tyr Tyr Arg Gly		2415
	2405	2410
Ser Asp Tyr Asp Glu Ala Asp Gly Pro Gly Ser Gly Gly Glu Glu		2430
2420	2425	2430
Ala Met Ala Gly Ala Tyr Asp Ala Pro Pro Pro Val Arg His Ala Ser		2445
2435	2440	2445
Ser Gly Ala Thr Gly Arg Ser Pro Arg Thr Pro Arg Ala Ser Gly Pro		2460
2450	2455	2460
Ala Cys Ala Ser Pro Ser Arg His Gly Arg Arg Leu Pro Asn Gly Tyr		2480
2465	2470	2475
Tyr Pro Ala His Gly Leu Ala Arg Pro Arg Gly Pro Gly Ser Arg Lys		2495
	2485	2490
Gly Leu His Glu Pro Tyr Ser Glu Ser Asp Asp Asp Trp Cys		2510
2500	2505	

<210> 11
 <211> 7791
 <212> DNA
 <213> Homo Sapiens

<400> 11	
gatgtcccga gctgctatcc ccggctcggc ccgggcagcc gccttctgag ccccgaccc	60
gaggcgccga gccgcgccg cccgatgggc tgggcctgag agcgtctccg cagtcgtagc	120
tccagccgcc gcgctcccag ccccgccagc ctcagcatca gcggcgccgc cgccggcgcc	180
ggcgtcttcc gcatcggttc ccgcagcgta acccgagacc ctttgcctct tgcagaatgg	240
cccgccttcg agacgagatg ccggcccgcct acgggggaggg aggcctccggg gcagccgccc	300
gggtgggtcgt gggcagcgga ggccggcgag gagccggggg cagccggcag ggccgggcagc	360
ccggggcgca aaggatgtac aagcagtcac tggcgccagag agcgccggacc atggcactct	420
acaaccccat ccccgctccga cagaactgcc tcacgggttaa ccgggtctctc ttccctcttca	480
gcgaagacaa cgtggtgaga aaatacgcca aaaagatcac cgaatggcct ccctttgaat	540
atatgatatt agccaccatc atagcgaatt gcacgtctct cgcactggag cagcatctgc	600
ctgatgatga caagaccccg atgtctgaac ggctggatga cacagaacca tacttcattg	660
gaattttttg tttcgaggct ggaattaaaa tcattgccct tgggtttgac ttccacaaag	720
gctcctactt gaggaatggc tgggaatgtca tggactttgt ggtggtgcta acgggcatct	780
tggcgacagt tgggacggag tttgacctac ggacgctgag ggcagttcga gtgctgccc	840
cgtcaagct ggtgtctgga atcccaagtt tacaagtcgt cctgaagtcg atcatgaagg	900
cgatgatccc tttgctgcag atcggcctcc tcctattttt tgcaatcctt atttttgcaa	960
tcatagggtt agaattttat atgggaaaat ttcataccac ctgctttgaa gaggggacag	1020
atgacattca gggtagtct ccggctccat gtgggacaga agagcccgcc cgcacctgcc	1080
ccaatgggac caaatgtcag cctactggg aagggcccaa caacgggatc actcagttcg	1140
acaacatcct gtttgcatg ctgactgttt tccagtgcac aaccatggaa ggggtggactg	1200
atctcctcta caatagcaac gatgcctcag ggaacacttg gaactggtt tacttcatcc	1260
ccctcatcat catcggtccc ttttttatgc tgaaccttgt gctgggtgtg ctgtcagggg	1320
agtttgccaa agaaagggaa cgggtggaga accggcgggc ttttctgaag ctgaggcgcc	1380
aacaacagat tgaacgtgag ctcaatgggt acatggaatg gatctcaaaa gcagaagagg	1440
tgatcctcgc cgaggatgaa actgacgggg agcagaggca tccctttgat ggagctctgc	1500
ggagaaccac cataaagaaa agcaagacag atttgctcaa ccccgagag gctgaggatc	1560

agctggctga	tatagcctct	gtgggttctc	ccttcgcccg	agccagcatt	aaaagtgcc	1620
agctggagaa	ctcgaccttt	tttcacaaaa	aggagaggag	gatgcgtttc	tacatccgcc	1680
gcatggtcaa	aactcaggcc	ttctactgga	ctgtactcag	tttggtagct	ctcaacacgc	1740
tgtgtgttgc	tattgtttcac	tacaaccagc	ccgagtggct	ctccgacttc	ctttactatg	1800
cagaattcat	tttcttagga	ctctttatgt	ccgaatggt	tataaaaatg	tacgggcttg	1860
ggacgcggcc	ttacttccac	tcttccttca	actgctttga	ctgtgggggt	atcattggga	1920
gcatcttcga	ggatcatctg	gctgtcataa	aacctggcac	atccttttga	atcagcgtgt	1980
tacgagccct	cagggttattg	cgtattttca	aagtcacaaa	gtactgggca	tctctcagaa	2040
acctggctgt	ctctctctct	aactccatga	agtccatcat	cagcctgttg	tttctccttt	2100
tcctgttcat	tgtctgtctt	gcccttttgg	gaatgcaact	cctcggcggc	cagtttaatt	2160
tcgatgaagg	gactcctccc	accaacttcg	atacttttcc	agcagcaata	atgacgggtg	2220
ttcagatcct	gacgggcgaa	gactggaacg	aggatcatgta	cgacgggata	aagtctcagg	2280
ggggcggtgca	ggggcgcatg	gtgtttctcca	tctattttcat	tgtactgacg	ctcttttggga	2340
actacacacct	cctgaatgtg	ttcttggcca	tcgtctggga	caatctggcc	aacgcccagg	2400
agctcaccaa	ggtggaggcg	gacgagcaag	aggaagaaga	agcagcgaa	cagaaacttg	2460
ccctacagaa	agccaaggag	gtggcagaag	tgagtcctct	gtccgcgcc	aacatgtcta	2520
tagctgtgaa	agagcaacag	aagaatcaaa	agccagccaa	gtccgtgttg	gagcagcgga	2580
ccagtgaagt	gcgaaagcag	aacttgctgg	cccgccggga	ggcctgtat	aacgaaatgg	2640
acccggacga	gcgctggaag	gctgcctaca	cgcggcacc	gcggccagac	atgaagacgc	2700
acttggaccg	gccgctggtg	gtggaccgcg	aggagaaccg	caacaacaac	accaacaaga	2760
gccgggcgcc	cgagcccacc	gtggaccagc	gcctcggcca	gcagcgcgcc	gaggacttcc	2820
tcaggaaaca	ggcccgcctac	cacgatcggg	ccggggacc	cagcggctcg	gcgggccttg	2880
acgcacggag	ggcctgggcg	ggaagccagg	agcccgagct	gagccgggag	ggaccctacg	2940
gccgcgagtc	ggaccaccac	gcccgggagg	gcagcctgga	gcaaccggg	ttctgggagg	3000
gcgaggccga	gcgaggcaag	gccggggacc	cccaccggag	gcacgtgcac	cggcaggggg	3060
gcagcaggga	gagccgcagc	gggtccccgc	gcacgggagc	ggacggggag	catcgacgtc	3120
atcgcgcgca	ccgcaggccc	ggggaggagg	gtccggaggga	caaggcgag	cggagggcgc	3180
ggcaccgcga	gggcagccgg	ccggcccggg	gcggcgaggg	cgagggcgag	ggccccgacg	3240
ggggcgagcg	caggagaagg	caccggcatg	gcgctccagc	cacgtacgag	ggggacgcgc	3300
ggagggagga	caaggagcgg	aggcatcgga	ggaggaaaga	gaaccagggc	tccggggctc	3360
ctgtgtcggg	ccccaacctg	tcaaccaccc	ggccaatcca	gcaggacctg	ggcccctaag	3420
acccaccctc	ggcagaggat	attgacaaca	tgaagaacaa	caagctggcc	accgcgaggt	3480
cgcccgctcc	ccacggcagc	cttggccacg	ccggcctgcc	ccagagccca	gccaagatgg	3540
gaaacagcag	cgacccccgc	cccatgctgg	ccatccctgc	catggccacc	aacccccaga	3600
acgcgcggag	ccgcgggacg	cccaacaaac	ccggggaacc	atccaatccc	ggccccccca	3660
agacccccga	gaatagcctt	atcgtcacca	acccagcg	caccagacc	aattcagcta	3720
agactgccag	gaaacccgac	cacaccacag	tggacatccc	cccagcctgc	ccaccccccc	3780
tcaaccacac	cgctgtacaa	gtgaacaaaa	acgccaaccc	agacccactg	ccaaaaaaag	3840
aggaagagaa	gaggaagacg	gaggtgggga	accgtgggga	agacggccct	aagccaaatg	3900
ctccctatag	ctccatgttc	atcctgtcca	cgaccaaccc	ccttcgccgc	ctgtgccatt	3960
acatcctgaa	cctgcgctac	tttgagatgt	gcacccctcat	ggtcattgcc	atgagcagca	4020
tcgcccctgg	cgccgaggac	cctgtgcagc	ccaacgcacc	tcggaacaac	gtgctgcgat	4080
acttttgcta	ggcgttctca	cctttgagat	cctttgagat	ggtgatcaag	atgattgacc	4140
tggggctcgt	cctgcatcag	ggtgcctact	tccgtgacct	ctggaatatt	ctcgacttca	4200
tagtggtcag	tggggccctg	gtagcctttg	ccttcaactgg	caatagcaaa	ggaaaagaca	4260
tcaacacgat	taaatccctc	cgagtccctc	gggtgctacg	acctcttaaa	accatcaagc	4320
ggctgccaaa	gctcaaggct	gtgtttgact	gtgtggtgaa	ctcacttaaa	aacgtcttca	4380
acatcctcat	cgtctacatg	ctattcatgt	tcactcttcg	cgtggtggct	gtgcagctct	4440
tcaaggggaa	attcttccac	tgcactgacg	agtccaaaga	gtttgagaaa	gattgtcgag	4500
gcaaatacct	cctctacgag	aagaatgagg	tgaaggcgcg	agaccgggag	tggagaagat	4560
atgaattcca	ttacgacaat	gtgctgtggg	ctctgctgac	cctcttcacc	gtgtccacgg	4620
gagaaggctg	gccacaggtc	ctcaagcatt	cgggtggacgc	cacctttgag	aaccagggcc	4680
ccagccccgg	gtaccgcatg	gagatgtcca	ttttctacgt	cgtctacttt	gtgggtgttc	4740
ccttctctct	tgtcaatatc	tttgtggcct	tgtatcatcat	caccttccag	gagcaagggg	4800
acaagatgat	ggaggaatac	agcctggaga	aaaatgagag	ggcctgcatt	gatttcgcca	4860
tcagcgccaa	gccgctgacc	cgacacatgc	cgcagaacaa	gcagagcttc	cagtaccgca	4920
tgtggcagtt	cgtgggtgtc	ccgcctttcg	agtacacgat	catggccatg	atcgccctca	4980
acaccatcgt	gcttatgatg	aagttctatg	gggcttctgt	tgcttatgaa	aatgcctctg	5040
gggtgttcaa	actcgtcttc	acctccctct	tctctctgga	atgtgtgctg	aaagtcatgg	5100
cttttgggat	tctgaattat	ttccgcgatg	cctggaacat	cttcgacttt	gtgactgttc	5160
tgggcagcat	caccgatatc	ctcgtgactg	agtttgggaa	tccgaataac	ttcatcaacc	5220
tgagctttct	cgcctctctc	cgagctgccc	ggctcatcaa	acttctccgt	cagggttaca	5280

```

ccatccgcat tcttctctgg acctttgtgc agtccttcaa ggcctgcct tatgtctgtc 5340
tgctgatcgc catgctcttc ttcattctatg ccatcattgg gatgcagggtg tttggtaaca 5400
ttggcatcga cgtggaggac gaggacagtg atgaagatga gttccaaatc actgagcaca 5460
ataacttccg gaccttcttc caggccctca tgcttctctt ccggagtgcc accggggaag 5520
cttggcacia catcatgctt tctgcctca gcggaacc gtgtgataag aactctggca 5580
tctgactcgc agagtgtggc aatgaatttg cttattttta ctttgttcc ttcattctcc 5640
tctgctcgtt tctgatgctg aatctctttg tcgcccgtcat catggacaac tttgagtacc 5700
tcacccgaga ctctcccatc ctgggcccc accacctgga tgagtacgtg cgtgtctggg 5760
ccgagtatga ccccgagct tggggccgca tgccttacct ggacatgtat cagatgtcga 5820
gacacatgtc tcacgcccctg ggtctgggga agaagtgtcc ggccagagtg gcttacaagc 5880
ggcttctcgc gatggacctg cccgtcgcag atgacaacac cgtccacttc aattccaccc 5940
tcatggctct gatccgcaca gccctggaca tcaagattgc caagggaagg gccgacaac 6000
agcagatgga cgctgagctg cggaaggaga tgatggcgat ttggcccaat ctgtcccaga 6060
agacgctaga cctgctgtgc acacctcaca agtccacgga cctcacctg ggaagatct 6120
acgcagccat gatgatcatg gagtactacc ggcagagcaa ggccaagaag ctgcaggcca 6180
tgcgcgaggga gcaggaccgg acacccctca tgttccagcg catggagccc ccgtcccaa 6240
cgcggaagg gggacctggc cagaacgccc tccccctcac ccagctggac ccaggaggag 6300
ccctgatggc tcacgaaaagc ggcctcaagg agaccgctc ctgggtgacc cagcgtcccc 6360
aggagatgtt ccagaagacg ggcacatgga gtccggaaca agggccccct accgacatgc 6420
ccaacagcca gcctaactct cagtccgtgg agatgcgaga gatgggcaga gatggctact 6480
ccgacagcga gcactacctc cccatggaag gccaggcgcc ggctgcctcc atgccccgcc 6540
tccctgcaga gaaccagagg agaaggggcc agccacgtgg gaataacctc agtaccatct 6600
cagacaccag ccccatgaag cgttcagcct ccgtgctggg cccaaggcc cgacgcctgg 6660
acgattactc gctggagcgg gtcccgcgcg aggagaacca gcggcaccac cagcgccgcc 6720
gcgaccgcag ccaccgcgcc tctgagcgct ccctggggcc ctacaccgat gtggacacag 6780
gcttggggac agacctgagc atgaccaccc aatccgggga cctgccgtcg aaggagcggg 6840
accaggagcg gggccggccc aaggatcgga agcatcgaca gcaccaccac caccaccacc 6900
accaccacca tccccgcgcc cccgacaagg accgctatgc ccaggaacgg ccggaccacg 6960
gccgggcacg ggctcgggac cagcgctggt cccgctcgcc cagcgagggc cgagagcaca 7020
tggcgaccgg gcagtagtgc cgtaagtgga agccagccc cctcaacatc tggtaaccg 7080
actccggggc gggggccgcc ccagctcccc cagacccccct ccaccccccg gccacacgtg 7140
tcctattccc ctgtgatccg taaggccggc ggctcggggc ccccgagca gcagcagcag 7200
cagcaggcgg tggccaggcc gggccggggc gccaccagcg gccctcgag gtaccaggc 7260
cccacggccg agcctctggc cggagatcgg ccgccacgg ggggccacag cagcgccgc 7320
tcgcccagga tggagaggcg ggtccpaggc ccggcccggg gcgagtcccc cagggcctgt 7380
cgacacggcg gggcccgggt gccggcatct ggcccgcacg tgtccgaggg gcccccgggt 7440
ccccggcacc atggctacta cgggggtccc gactacgacg agggcgatgg cccgggcagc 7500
ggggcgggcg aggagccat ggccggggcc tacgacgcgc cacccccct acgacacgcg 7560
tctcggggcg ccaccggggc ctgcccagg actccccgg cctcggggcc ggctgcgcc 7620
tcgcttctc ggcacggccg gcgactccc aacggctact accggcgca cggactggcc 7680
aggccccgcg ggccgggctc cagggaaggc ctgcacgaac cctacagcga gagtgcagat 7740
gattggtgct aagcccgggc gaggtggcgc ccgcccggcc cccacgcac c 7791

```

<210> 12

<211> 2266

<212> PRT

<213> Homo Sapiens

<400> 12

```

Met Ala Arg Phe Gly Asp Glu Met Pro Ala Arg Tyr Gly Gly Gly Gly
1      5      10      15
Ser Gly Ala Ala Ala Gly Val Val Val Gly Ser Gly Gly Gly Arg Gly
20     25     30
Ala Gly Gly Ser Arg Gln Gly Gly Gln Pro Gly Ala Gln Arg Met Tyr
35     40     45
Lys Gln Ser Met Ala Gln Arg Ala Arg Thr Met Ala Leu Tyr Asn Pro
50     55     60
Ile Pro Val Arg Gln Asn Cys Leu Thr Val Asn Arg Ser Leu Phe Leu
65     70     75     80
Phe Ser Glu Asp Asn Val Val Arg Lys Tyr Ala Lys Lys Ile Thr Glu

```

- 38 -

- 39 -

- 40 -

1570	1575	1580
Val Leu Met Met Lys Phe Tyr Gly Ala Ser	Val Ala Tyr Glu Asn Ala	
1585	1590	1595
Leu Arg Val Phe Asn Ile Val Phe Thr Ser Leu Phe Ser Leu Glu Cys		1600
	1605	1610
Val Leu Lys Val Met Ala Phe Gly Ile Leu Asn Tyr Phe Arg Asp Ala		1615
	1620	1625
Trp Asn Ile Phe Asp Phe Val Thr Val Leu Gly Ser Ile Thr Asp Ile		1630
	1635	1640
Leu Val Thr Glu Phe Gly Asn Pro Asn Asn Phe Ile Asn Leu Ser Phe		1645
	1650	1655
Leu Arg Leu Phe Arg Ala Ala Arg Leu Ile Lys Leu Leu Arg Gln Gly		1660
1665	1670	1675
Tyr Thr Ile Arg Ile Leu Leu Trp Thr Phe Val Gln Ser Phe Lys Ala		1680
	1685	1690
Leu Pro Tyr Val Cys Leu Leu Ile Ala Met Leu Phe Phe Ile Tyr Ala		1695
	1700	1705
Ile Ile Gly Met Gln Val Phe Gly Asn Ile Gly Ile Asp Val Glu Asp		1710
	1715	1720
Glu Asp Ser Asp Glu Asp Glu Phe Gln Ile Thr Glu His Asn Asn Phe		1725
	1730	1735
Arg Thr Phe Phe Gln Ala Leu Met Leu Leu Phe Arg Ser Ala Thr Gly		1740
1745	1750	1755
Glu Ala Trp His Asn Ile Met Leu Ser Cys Leu Ser Gly Lys Pro Cys		1760
	1765	1770
Asp Lys Asn Ser Gly Ile Leu Thr Arg Glu Cys Gly Asn Glu Phe Ala		1775
	1780	1785
Tyr Phe Tyr Phe Val Ser Phe Ile Phe Leu Cys Ser Phe Leu Met Leu		1790
	1795	1800
Asn Leu Phe Val Ala Val Ile Met Asp Asn Phe Glu Tyr Leu Thr Arg		1805
	1810	1815
Asp Ser Ser Ile Leu Gly Pro His His Leu Asp Glu Tyr Val Arg Val		1820
1825	1830	1835
Trp Ala Glu Tyr Asp Pro Ala Ala Trp Gly Arg Met Pro Tyr Leu Asp		1840
	1845	1850
Met Tyr Gln Met Leu Arg His Met Ser Pro Pro Leu Gly Leu Gly Lys		1855
	1860	1865
Lys Cys Pro Ala Arg Val Ala Tyr Lys Arg Leu Leu Arg Met Asp Leu		1870
	1875	1880
Pro Val Ala Asp Asp Asn Thr Val His Phe Asn Ser Thr Leu Met Ala		1885
	1890	1895
Leu Ile Arg Thr Ala Leu Asp Ile Lys Ile Ala Lys Gly Gly Ala Asp		1900
1905	1910	1915
Lys Gln Gln Met Asp Ala Glu Leu Arg Lys Glu Met Met Ala Ile Trp		1920
	1925	1930
Pro Asn Leu Ser Gln Lys Thr Leu Asp Leu Leu Val Thr Pro His Lys		1935
	1940	1945
Ser Thr Asp Leu Thr Val Gly Lys Ile Tyr Ala Ala Met Met Ile Met		1950
	1955	1960
Glu Tyr Tyr Arg Gln Ser Lys Ala Lys Lys Leu Gln Ala Met Arg Glu		1965
	1970	1975
Glu Gln Asp Arg Thr Pro Leu Met Phe Gln Arg Met Glu Pro Pro Ser		1980
1985	1990	1995
Pro Thr Gln Glu Gly Gly Pro Gly Gln Asn Ala Leu Pro Ser Thr Gln		2000
	2005	2010
Leu Asp Pro Gly Gly Ala Leu Met Ala His Glu Ser Gly Leu Lys Glu		2015
	2020	2025
Ser Pro Ser Trp Val Thr Gln Arg Ala Gln Glu Met Phe Gln Lys Thr		2030
	2035	2040
Gly Thr Trp Ser Pro Glu Gln Gly Pro Pro Thr Asp Met Pro Asn Ser		2045
	2050	2055
Gln Pro Asn Ser Gln Ser Val Glu Met Arg Glu Met Gly Arg Asp Gly		2060

2065						2070						2075				2080
Tyr	Ser	Asp	Ser	Glu	His	Tyr	Leu	Pro	Met	Glu	Gly	Gln	Gly	Arg	Ala	
				2085					2090						2095	
Ala	Ser	Met	Pro	Arg	Leu	Pro	Ala	Glu	Asn	Gln	Arg	Arg	Arg	Gly	Arg	
			2100					2105						2110		
Pro	Arg	Gly	Asn	Asn	Leu	Ser	Thr	Ile	Ser	Asp	Thr	Ser	Pro	Met	Lys	
		2115					2120					2125				
Arg	Ser	Ala	Ser	Val	Leu	Gly	Pro	Lys	Ala	Arg	Arg	Leu	Asp	Asp	Tyr	
	2130					2135					2140					
Ser	Leu	Glu	Arg	Val	Pro	Pro	Glu	Glu	Asn	Gln	Arg	His	His	Gln	Arg	
2145					2150				2155						2160	
Arg	Arg	Asp	Arg	Ser	His	Arg	Ala	Ser	Glu	Arg	Ser	Leu	Gly	Arg	Tyr	
				2165					2170					2175		
Thr	Asp	Val	Asp	Thr	Gly	Leu	Gly	Thr	Asp	Leu	Ser	Met	Thr	Thr	Gln	
			2180					2185					2190			
Ser	Gly	Asp	Leu	Pro	Ser	Lys	Glu	Arg	Asp	Gln	Glu	Arg	Gly	Arg	Pro	
	2195						2200					2205				
Lys	Asp	Arg	Lys	His	Arg	Gln	His	His	His	His	His	His	His	His	His	
	2210					2215					2220					
His	Pro	Pro	Pro	Pro	Asp	Lys	Asp	Arg	Tyr	Ala	Gln	Glu	Arg	Pro	Asp	
2225					2230				2235					2240		
His	Gly	Arg	Ala	Arg	Ala	Arg	Asp	Gln	Arg	Trp	Ser	Arg	Ser	Pro	Ser	
			2245					2250						2255		
Glu	Gly	Arg	Glu	His	Met	Ala	His	Arg	Gln							
			2260					2265								

(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property
Organization
International Bureau



(43) International Publication Date
23 January 2003 (23.01.2003)

PCT

(10) International Publication Number
WO 2003/006103 A3

(51) International Patent Classification⁷: **C07H 21/04**,
C12N 5/00, 15/00, C12P 1/06, G01N 33/53, 33/554,
35/00, G01R 27/00

(US). MCMANUS, Owen, B. [US/US]; 34 Robin Drive,
Skillman, NJ 08558 (US).

(21) International Application Number:
PCT/US2002/022161

(74) Agent: VAN DYKE, Timothy, H.; Van Dyke & Asso-
ciates, P.A., 1630 Hillcrest Street, Orlando, FL 32803 (US).

(22) International Filing Date: 12 July 2002 (12.07.2002)

(81) Designated States (*national*): AE, AG, AL, AM, AT, AU,
AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU,
CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH,
GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC,
LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW,
MX, MZ, NO, NZ, OM, PH, PL, PT, RO, RU, SD, SE, SG,
SI, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ,
VN, YU, ZA, ZM, ZW.

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Data:
60/304,955 12 July 2001 (12.07.2001) US

(84) Designated States (*regional*): ARIPO patent (GH, GM,
KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW),
Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM),
European patent (AT, BE, BG, CH, CY, CZ, DE, DK, EE,
ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, SK,
TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GQ,
GW, ML, MR, NE, SN, TD, TG).

(71) Applicant (*for all designated States except US*): MERCK
& CO., INC. [US/US]; Patent department, P.O. Box 2000
- RY60-30, Rahway, NJ 07065-0907 (US).

(72) Inventors; and

(75) Inventors/Applicants (*for US only*): AUGUSTINE,
Paul, R. [US/US]; 8 Stima Avenue, Carteret, NJ 07008
(US). BENNETT, Paul, B. [US/US]; 3679 Hancock Lane,
Doylestown, PA 18901 (US). BUGIANESI, Randal, M.
[US/US]; 475 Milcrip Road, Bridgewater, NJ 09907 (US).
GARYANTES, Tina, A. [US/US]; 18 Roberts Road,
Warren, NJ 07059 (US). IMREDY, John, P. [US/US]; 861
Yorktown Street, Lansdale, PA 19446 (US). KATH, Gary,
S. [US/US]; 2671 Sky Top Drive, Scotch Plains, NJ 07076

Published:

— with international search report

(88) Date of publication of the international search report:
25 March 2004

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

(54) Title: ELECTRICAL FIELD STIMULATION OF EUKARYOTIC CELLS

(57) Abstract: Methods of identifying activators and inhibitors of voltage-gated ion channels are provided in which the methods employ electrical field stimulation of the cells in order to manipulate the open/close state transition of the voltage-gated ion channels. This allows for more convenient, more precise experimental manipulation of these transitions, and, coupled with efficient methods of detecting the result of ion flux through the channels, provides methods that are especially suitable for high throughput screening.

WO 2003/006103 A3

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US02/22161

A. CLASSIFICATION OF SUBJECT MATTER

IPC(7) : C07H 21/04; C12N 5/00, 15/00; C12P 1/06; G01N 33/53, 33/554, 35/00; G01R 27/00
 US CL : 324/600; 422/50, 55, 67; 435/69.1, 320.1, 325; 436/43, 519, 800, 807, 809; 536/23.5

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

U.S. : 324/600; 422/50, 55, 67; 435/69.1, 320.1, 325; 436/43, 519, 800, 807, 809; 536/23.5

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)
 USPT, PGPB, JPAB, EPAB, DWPI, REGISTRY, HCAPLUS

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Y	US 6,057,114 A (AKONG et al.) 02 May 2000 (02.05.2000), Abstract, Column 1, Lines 12-15; Column 4, Lines 4-22; Column 9, Lines 63-68; Column 17, Lines 22-40; Column 17, Line 63 to Column 18, Line 15; Column 20, Lines 32-54; Column 22, Line 57 to Column 23, Line 30; Column 23, Line 54 to Column 24, Line 9; Column 26, Lines 9-20; Column 27, Lines 14-22; Column 33, Line 61 to Column 34 Line 27; Column 41, Line 43 to Column 42, Line 9; Column 42, Lines 28-67; Column 43, Lines 35-56.	1-16 and 20-60
Y	CONNOLLY, P. et al. An Extracellular Microelectrode Array for Monitoring Electrogenic Cells in Culture. Biosensors and Bioelectronics, 1990, Vol. 5, Pages 223-	17-19 and 61-74
Y, P	US 6,377,057 B1 (BORKHOLDER) 23 April 2002 (23.04.2002), entire document.	17-19 and 61-74

☐ Further documents are listed in the continuation of Box C.☐ See patent family annex.

* Special categories of cited documents:

"A" document defining the general state of the art which is not considered to be of particular relevance

"E" earlier application or patent published on or after the international filing date

"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)

"O" document referring to an oral disclosure, use, exhibition or other means

"P" document published prior to the international filing date but later than the priority date claimed

"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art

"Δ" document member of the same patent family

Date of the actual completion of the international search

27 August 2003 (27.08.2003)

Date of mailing of the international search report

11 DEC 2003

Name and mailing address of the ISA/US

Mail Stop PCT, Attn: ISA/US
 Commissioner for Patents
 P.O. Box 1450
 Alexandria, Virginia 22313-1450
 Facsimile No. (703)305-3230

Authorized officer.

Dr. Kailash C. Srivastava

Telephone No. (703)-308-0196

**This Page is Inserted by IFW Indexing and Scanning
Operations and is not part of the Official Record**

BEST AVAILABLE IMAGES

Defective images within this document are accurate representations of the original documents submitted by the applicant.

Defects in the images include but are not limited to the items checked:

- ☐ BLACK BORDERS
- ☐ IMAGE CUT OFF AT TOP, BOTTOM OR SIDES
- ☐ FADED TEXT OR DRAWING
- ☐ BLURRED OR ILLEGIBLE TEXT OR DRAWING
- ☐ SKEWED/SLANTED IMAGES
- ☒ COLOR OR BLACK AND WHITE PHOTOGRAPHS
- ☐ GRAY SCALE DOCUMENTS
- ☒ LINES OR MARKS ON ORIGINAL DOCUMENT
- ☐ REFERENCE(S) OR EXHIBIT(S) SUBMITTED ARE POOR QUALITY
- ☐ OTHER: _____

IMAGES ARE BEST AVAILABLE COPY.

As rescanning these documents will not correct the image problems checked, please do not report these problems to the IFW Image Problem Mailbox.